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is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
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## ALIGNMENTS

RESULT 1 US-08-875-233-5

Sequence 5, Application US/08875233
Patent No. 6127601
GENERAL INFORMATION:

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US-08-875-233-5
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Patent No.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 312-616-5460
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/875,233 FILING DATE: 26-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Boeshore, Maury L
APPLICANT: McMaster, J. Russell
APPLICANT: Tricoli, David M
APPLICANT: Reynolds, John F
APPLICANT: Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C
TITLE OF INVENTION: Cucumber Mosaic Virus
NUMBER OF SEQUENCES: 15
                                                                                                                                                        FEATURE:
NAME/KEY:
                                                                                                                                                                                      ORIGINAL SOURCE:
ORGANISM: Cucumber Mosaic Virus
INDIVIDUAL ISOLATE: A35
                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: 312-616-5400
                                                                                                                                                                                                                                        MOLECULE TYPE:
HYPOTHETICAL: NO
ANTI-SENSE: NO
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                                                                    Local Similarity
                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                        LOCATION:
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; Sequence 3, Application US/08875233; Patent No. 6127601; GENERAL INFORMATION: APPLICANT: Boeshore, Maury L APPLICANT: McMaster, J. Russell APPLICANT: Tricoli, David M APPLICANT: Reynolds, John F
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INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 792 base pairs
TYPE: nucleic acid
                                                                                                                                                         Patent No. 5641673
                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Brand, Andrea
APPLICANT: Perrimon, No. 5641673bert
APPLICANT: Goodman, Howard M.
TITLE OP INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                              APPLICANT: Haseloff, James
APPLICANT: Brand, Andrea
APPLICANT: Perrimon, No. 5
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MEDIUM TYPE: Floppy
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TITLE OF INVENTION: Plants Resistant to C Strains
TITLE OF INVENTION: Cucumber Mosaic Virus
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Local Similarity 100.0%; Pred. No. 5.9e-:
nes 46; Conservative 0; Mismatches
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LOCATION: 3..660
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FILING DATE: 26-JUN-1997
CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN: v-33
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Y: U.S.A.
                                                                                                                                                                             Application US/08090193
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CORRESPONDENCE ADDRESS:

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US-08-488-031-14
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PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/642,330
APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
NAME: Cimbala, Michele A.
NAME: 33,851
                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 14,
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                                                                                                                                                                                                                                                                                                                                                                                                   Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                              APPLICANT: Haseror,
APPLICANT: Brand, Andrea
APPLICANT: Berrimon, No. 5849548bert
APPLICANT: Goodman, Howard M.
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
TITLE OF TOTELES: 56
                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA
APPLICATION NUMBER: US/08/488,031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: U$/08/090,193
FILING DATE: 23-DEC-1993
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PRIOR APPLICATION DATA:
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CITY: Washington
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STRANDEDNESS: both
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FILING DATE: 07-JUN-
IOR APPLICATION DATA:
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                  07-JUN-1995
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100.0%; Pred. No. 2.3e-1
tive 0; Mismatches
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APPLICANT: Haselo
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                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: PC-DOS/MS-DOS
                                                                                                                                                                PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: PCT/US92/00277
                                                                                                                                                                                                             APPLICATION NUMBER: US/08/486,569
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,193
FILING DATE: 23-DEC-1993
                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
THE //OF/ARK 569
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LENGTH: 130 base pairs
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080006/MAC/LBB
TELECOMMUNICATION INFORMATION:
                                                                                                                                    APPLICATION NUMBER: PCT/I
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 5
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ATTORNEY/AGENT INFORMATION:
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                                                                         FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
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TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
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STRANDEDNESS: both
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REFERENCE/DOCKET NUMBER: 0609.3080002/MAC/LBB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Bugaisky, Lawrence B. REGISTRATION NUMBER: 35,086
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                                    NAME: Bugaisky, Lawrence B
REGISTRATION NUMBER: 35,08
                                                                                               APPLICATION NUMBER: US 07/642,330 FILING DATE: 17-JAN-1991
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Perrimon, No. 5863774bert
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100.0%; Pred. No. 2.3e-15;
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RESULT 7
US-08-488-027-14
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                                                                                                              US-08-488-027-14
                                    Matches
                                                      Query Match
Best Local Similarity
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PRIOR APPLICATION DATA: 08/090,193
APPLICATION NUMBER: 08/090,193
PRIOR APPLICATION DATA: 23-DEC-1993
PRIOR APPLICATION DATA: PCT/US92/00277
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
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LENGTH: 130 base pairs
                                                                                                                                                                                                                                                                            NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080003/MAC/LBB
TELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: US (
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
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TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                            LENGTH: 130 base pairs TYPE: nucleic acid STRANDEDNESS: both TOPOLOGY: linear
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FILING DATE: 16-JAN-1992
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ZIP: 20005
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                                    45; Conservative
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5 CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
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1100 New York Avenue, N.W., Suite 600
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Perrimon, No. 5866384bert
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 84.9%; bu-
100.0%; Pred. No.
100.0%; Mismatches
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100.0%; Pred. No.
tive 0; Mismatc
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                                 Score 45; DB 2; Length 130; Pred. No. 2.3e-15; O; Mismatches 0; Indels
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US-08-090-192-14
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                                                                                                                                                                                                                                US-08-482-663-14
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                                                                                                                                                                                        Sequence 14, Application US/08482663 Patent No. 5882907
                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
                             GENERAL INFORMATION:
APPLICANT: Haseloff, James
APPLICANT: Haseloff, James
APPLICANT: Brand, Andrea
APPLICANT: Perrimon, No. 5882907bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
NUMBER OF SEQUENCES ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (202) 371-25
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0609
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 130 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PCT/ US 92/00278
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 11-NOV-
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Haseloff, APPLICANT: Goodman, I
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STRANDEDNESS: b
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
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E: Sterne, Kessler, Goldstein & Fox 1100 New York Avenue, N.W., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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100.0%; Pred. No. 2.3e-15;
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3ER: 0609.3030001
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COMPUTER READABLE FORM: MEDIUM TYPE: Floppy

Floppy disk

20005

STATE: C COUNTRY:

D.C.

U.S.A.

Washington

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RESULT 10
US-08-482-658-14
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SEQUENCE CHARACTERISTICS:
TENGTH: 130 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 14, Application US/08482658 Patent No. 6010904
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CURENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,663
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,193
FILING DATE: 23-DEC-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Haselo
                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,658
                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                               APPLICANT: Brand, Andrea
APPLICANT: Perrimon, No. 6010904bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 07/0
PILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
TELEPAX: (202) 371-2540
                   FILING DATE: 07-JUN-1995 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                      STREET: 1100 New
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                        COUNTRY:
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OPERATING SYSTEM: PC-DOS/MS-DOS
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    Sterne, Kessler, Goldstein & Fox
    1100 New York Avenue, N.W., Suite 600

                                                                                                                                                                                                          U.S.A.
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100.0%; Pred. No.
08/090,193
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US-08-470-349-14
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                                                                                                                                                                          CLASSIFICATION: 435
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,192
APPLICATION NUMBER: PCT/ US 92/00278
APPLICATION NUMBER: PCT/ US 92/00278
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 14, Application US/08470349 Patent No. 6015794
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (202) 371-254
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US
FILING DATE: 16-CAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin NDATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,349
FILING DATE: 06-UN-1995
FILING DATE: 06-UN-1995
                                                                                                                             APPLICATION NUMBER: US 07 FILING DATE: 17-JAN-1991 ATTORNEY/AGENT INFORMATION: NAME: Cimbala, Michele A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                       REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Haselott, James
APPLICANT: Goodman, Howard M.
APPLICANT: Trans-Splicing Ribozymes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 5
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                                                                                                                 REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Bugaisky, Lawrence B. REGISTRATION NUMBER: 35,086
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                                                       TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Washington
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X: (202) 371-2540
N FOR SEQ ID NO:
CHARACTERISTICS:
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(202) 371-2540
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                                                         (202)
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                                                       371-2600
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100.0%; Pred. No. 2.3e-15;
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                                                                                                             33,851
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                                                                                               0609.3030001
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RESULT 12
US-08-475-610-14
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/475,610
ETLING DATE: 07-UN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,193
ETLING DATE: 23-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JRAT-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JRAT-1992
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/642,330
PTLING DATE: 17-JRAT-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 14, Application US/08475610 Patent No. 6071730
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Best Local Similarity
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                                                                                                                             Query Match
                                                                                                                                                                                                                                                                           NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080004/MAC/LBB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
TELEPAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox
STREET: 1100 New York Avenue, N.W., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Perrimon, No. 6071730bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 130 base pairs TYPE: nucleic acid STRANDEDNESS: both
                                                                                                                                                                                                                 LENGTH: 130 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                           Local
                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tch 84.9%; Score 45; al Similarity 100.0%; Pred. No. 45; Conservative 0; Mismatci
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                                                                                     Similarity 45; Conserv
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                             CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
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                                                                                   84.9%; Score 45; DB 3; Lilarity 100.0%; Pred. No. 2.3e-15; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                 linear
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                                                                                                                             Length 130,
                                                                                       Indels
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PCT-US92-00278-14
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PCT-US92-00277-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCT-US92-00277-14
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
ZIP: 20036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Goldstein, Jorge A
REGISTRATION NUMBER: 29,021
REFERENCE/DOCKET NUMBER: 0609.3496604
TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: 07/454517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CURRENT APPLICATION DATA:
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APPLICANT: Perrimon, Norbert
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                        STREET: La..
STREET: Washington
CITY: Washington
CTATE: DC
TTATE: TTATE
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APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: NUCLEIC ACID STRANDEDNESS: both
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                                                                                                                                                                                                    COUNTRY:
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1225 Connecticut Avenue, N.W., Suite 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            130 bases
                                                                                                                                                                                                                                                                                                                                    E: Sterne, Kessler, Goldstein & Fox 1225 Connecticut Avenue, N.W., Suit
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 84.9%; Score 45; DB 5; 100.0%; Pred. No. 2.3e-1
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US-08-090-193-11
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SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/090,193
FILING DATE: 23-DEC-1993
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UNUMBER: PCT/US92/00277
FILING DATE: 16-JAN-1992
PRIOR APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
REGISTRATION NUMBER: 33.0851
REFERENCE/DOCKET NUMBER: 0609.3080001
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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Best Local S
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TELEFAX: (202)833-8716
INFORMATION FOR SEQ ID NO: 14
SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
PRIOR APPLICATION UNMBER: US 07/642,333
PILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jozge A
REGISTRATION NUMBER: 29,021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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REFERENCE/DOCKET NUMBER: 0609.3476604
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox
STREET: 1100 New York Avenue, Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: PC-D
SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                  20005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         T: Brand, Andrea
T: Perrimon, No. 5641673bert
T: Goodman, Howard M.
INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            130 bases
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In Release #1.0, Version #1.25
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hes 0;
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Query Match
Best Local Similarity
Watches 45; Conserva
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                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                              LENGTH: 131 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: both
                                            53 CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 97
                                                            5 CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                                                       Conservative
                                                                                                    84.9%; Score 45; DB 1; Le
100.0%; Pred. No. 2.3e-15;
n. Mismatches 0;
5, 2004, 08:00:07
                                                                                                                                 Length 131,
                                                                                                       Indels
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ABM637180
AAC21510
AAC21509
ADA731508
ABM86674
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AAR67753
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Pseudomor	Human sec	Propionib	Propionib	Human dig	Pseudomon	HSV-2 str	Human sec	Pseudomon	Human imm	Propionib	Propionib	Peptide	~	MHC/HLA p					

ALIGNMENTS

AAY90255 standard; protein; 14 A

AAY90255;

**Ubiquitin** monomer C-terminal 19-SEP-2000 (first entry) fragment

Ubiquitin monomer; protein production; plant cell; ubiquitin promoter.

Nicotiana tabacum

WO200036129-A1.

22-JUN-2000.

11-DEC-1998; 98WO-SG006103

(MOLE-) INST MOLECULAR AGROBIOLOGY.

11-DEC-1998;

98WO-SG000103

Fang R, Wu J, Chen

WPI; 2000-431604/37.

N-PSDB; AAA30858

Production of desired protein in plants or plant cells ubiquitin monomer coding sequence upstream of the gene desired protein. by linking a encoding the

Claim 7; Page 18; 42pp; English.

This sequence represents the C-terminal fragment of a ubiquitin monomer The invention relates to a method for enhancing production of a desired protein in a plant or plant cell by inserting a nucleic acid (NA) encoding a ubiquitin monomer upstream of a NA encoding the desired protein, where the fusion construct encodes a fusion protein and expression is not controlled by the ubiquitin promoter. The invention also relates to a NA acid vector a NA vector able to transform a plant cell, that comprises NA encoding a fusion protein having a ubiquitin monomer linked to a protein of interest and further, where expression of the fusion construct is not under control of a ubiquitin promoter. The construct allows enhanced production of the desired protein in plants of the construct allows enhanced production of the desired protein in plants of the construct allows enhanced production of the desired protein in plants of the construct allows enhanced production of the desired protein in plants of the construct allows enhanced production of the desired protein in plants of the construct allows enhanced production of the desired protein in plants of the construct allows enhanced production of the desired protein in plants of the construct allows enhanced production of the desired protein in plants of the construct allows enhanced production of the desired protein in plants of the construct allows enhanced production of the desired production in plants. desired õ of.

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RESULT 2
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XX EP41
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XX I1-;
AR HIC
XX CAP5
XX CAP5
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RESULT 3
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Best Local :
                                                                                                                                                                                                                                                                                      The gene encoding this protein was isolated from a plasmid (pUC18) containing DNA complementary to RNA 3 of the virulent strain FNY, isolated in New York on infected melons. Transgenic melon plantlets containing the nucleotide sequence introduced via an Agrobacterium tumefaciens intermediate are cultured as shoots in special media. The transformed melon plants are resistant to CMV. See also AAQ10462. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 correct PI field.) (Updated on 27-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                            Sequence 218 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Transgenic melon plantlets prodn. from transformed shoots - by two-stage culture on specific medium for new transformed plants etc., esp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
      AAR10653;
                                      AAR10653 standard; protein; 218
                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 15; Page 17; 44pp; French.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CMV; resistance; capsid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Capsid protein of Cucumber Mosaic Virus strain FNY.
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25-MAR-2003
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                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                          Score 14; DB 2;
Pred. No. 1.7e-06;
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Pred. No. 1.6e-07;
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                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                          0
                                                                                                                                                                                                                         Length 218;
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25-MAR-2003
16-APR-1991
                        26-FEB-1993;
                                                         02-SEP-1994.
                                                                                                                                                         polyribozyme;
                                                                                                                                                                                                          Cauliflower mosaic virus capsid protein.
                                                                                                                                                                                                                                           25-MAR-2003
20-APR-1995
                                                                                                                                                                                                                                                                                             AAR57968;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 16; Page 19; 44pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    resistant to cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Deboth M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-AUG-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-AUG-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EP412912-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cucumber mosaic
                                                                                                                         Cauliflower mosaic
                                                                                                                                                                                                                                                                                                                              AAR57968 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-FEB-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BIOC-) BIOCEM SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pocar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1991-046027/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                              1 MDKSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                MDKSESTSAGRNRR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 218 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                           (revised)
(first entry)
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                        93FR-00002269
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          virus
                                                                                                                                                                                                                                                                                                                           protein; 218
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The gene encoding this protein was isolated from the virulent French strain 117F of CMV. Transgenic melon plantlets containing the coding sequence (introduced via an Agrobacterium tumefaciens intermediate) are cultured as shoots in special media. The transformed melon plants are resistant to CMV. See also AAQ10461. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PI field.) (Updated on 25-MAR-2003 to correct PI field.)
Cauliflower mosaic virus; CMV; capsid protein; coat protein; polyribozyme; inactivate; inactivation; resistance; crop pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Transgenic melon plantlets prodn. from transformed shoots - culture on specific medium for new transformed plants etc.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CMV; strain 117F; resistance; capsid protein; Cucumis melo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Capsid protein of Cucumber Mosaic Virus strain 117F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14; DB 2;
Pred. No. 1.7e-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.7e-06;
hes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 218;
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protection
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>.</u>
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RESULT 5
AAR67753
ID AAR6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New polyribozyme contg. several catalytic regions in complementary sequence - can inactivate gene for viral capsid protein, esp. for pre of new virus resistant transgenic plants, also DNA sequence encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
DNA constructs for imparting pathogen, encodes RNA that interacts with viral reliciting agent.
                                                                                                                                                                                                                                                                  virus resistance; disease resistance; transgenic plant; hypersensitive response; crop improvement; tomato; Lycopersicon esculentum; tobacco; Nicotiana tabacum; coa
                                                                                                                                            04-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
20-JUL-1995
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                                                                             De Haan
                                                                                                             (SANO)
                                                                                                                                                                  03-JUN-1994;
                                                                                                                                                                                        22-DEC-1994.
                                                                                                                                                                                                              WO9429464-A1
                                                                                                                                                                                                                                    Cucumber mosaic
                                                                                                                                                                                                                                                                                                   Cucumber mosaic virus; tomato mosaic virus; minus-strand RNA;
                                                                                                                                                                                                                                                                                                                          CMV coat protein
                                                                                                                                                                                                                                                                                                                                                                                                       AAR67753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 218
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                                                                                                   (SANO)
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                                             1995-036490/05.
DB; AAQ76106.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14;
                                                                                                 SANDOZ LTD.
SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN VERW GES MBH
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                                                                                                                                                                                                                                                                                                                                                                                                       standard;
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(first en
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                                                                                                                                                                  94WO-EP001817
                                                                                                                                                                                                                                                                                                                                                                                                     protein;
                                                                                                                                                                                                                                                                                                                                                entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14;
Pred. No.
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           RNA
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J. 1.7e-06;
O;
          esp. virus, resistance to RNA polymerase to generate
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                                                                                                                                                                                                                                                                                        cross protection;
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MDKSESTSAGRNRR

Query Match Best Local (

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218 AA;

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Similarity

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Score 14; 1 Pred. No. 1

DB 2; 1 1.7e-06; les 0;

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RESULT (
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ID AAR93803

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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     coat protein (CP) of tomato mosaic virus (AAR67552) as well as its own CP (AAR67753). The construct elicits minus-sense RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus resistance on a host plant, e.g. tobacco, tomato. (Updated on 25-MAR-2003 to correct PN field.)
                                              Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucum mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers backnown CP sequences. The genes may be used for producing plants such quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                         New isolated cucumber mosaic virus coat protein plants, partic. of the family Cucurbitaceae or sresistant to infection.
                                                                                                                                                                                                                      Disclosure; Fig 8; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-333993/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cucumber mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Coat protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-JUL-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of the A35 strain of cucumber mosaic virus
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(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ÇP;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 2; L.
1.7e-06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reynolds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CMV; V27;
                                                                                                                                                                                                                                                                                                n DNA - used
Solanaceae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 218;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     JF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             V33; V34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Carney
                                                                                                                                                                                                                                                                                                   to produce which are
                                                                                                                                                                           of cucumber
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RESULT 7
AAR99893
ID AAR9
AX AAR9
AX AAR9
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AX Coat
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RESULT 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based oknown CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae, resistant to infection.
                                                                                                                                                                                        16-OCT-2003
02-DEC-1996
                                                                                                                                                                                                                                                                   AAR98894;
               Cucumber mosaic
                                                                                     Coat protein; CP;
                                                                                                                                       Coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Fig 1; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-333993/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Boeshore ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  W09621018-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cucumber mosaic virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Coat protein; CP; cucumber mosaic virus; CMV; V27; V33; V34; A35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Coat protein of the V27 strain of cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-OCT-2003
02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR98893;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR98893 standard; protein; 218
                                                                                                                                                                                                                                                                                                               AAR98894 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ASGR-) ASGROW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                               μ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MDKSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                         MDKSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MDKSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA;
                                                                                                                                    of the V33 strain of cucumber mosaic virus.
                                                                                                                                                                                     (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mcmaster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94US-00367789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95WO-US007234
            virus; strain V33
                                                                                        cucumber mosaic virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; 8
100.0%; 8
tive 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ďŔ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 strain V27.
                                                                                                                                                                                                                                                                                                                  218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tricoli DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14; DB 2;
Pred. No. 1.7e-06;
Mismatches 0;
                                                                                                                                                                                                                                                                                                                  B
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                                                                                        CMV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Reynolds JF, Carney KJ;
                                                                                        V27; V33; V34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 218;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               to produce which are
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RESULT 9
AAR98895
ID AAR9
AC AAR9
AC AAR9
DT 16-C
DT 02-D
XX Coat
XX Coat
XX Coat
XX Coat
XX Coat
XX Coat
XX OAS
DR W096
PN W096
PN W096
XX W11-J
XX J
PR 07-J
XX I
PR 07-J
XX J
PR 07-J
XX I
PR 07-J
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                       WPI; 199
N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-OCT-2003
02-DEC-1996
                                                                                                                                                                                                                                    30-DEC-1994;
                                                                                                                                                                                                                                                                               07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Coat protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR98895 standard; protein; 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Fig 2; 80pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB;
                                                                                                                                            Boeshore ML,
                                                                                                                                                                                                                                                                                                                            11-JUL-1996.
                                                                                                                                                                                                                                                                                                                                                                         WO9621018-A1
                                                                                                                                                                                                                                                                                                                                                                                                                    Cucumber mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                               resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR98895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1996-333993/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Boeshore ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-JUL-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   resistant to infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9621018-A1
                                                                                                                                                                                       (ASGR-) ASGROW SEED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                       1996-333993/33.
DB; AAT34666.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 MDKSESTSAGRNRR 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ASGROW SEED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MDKSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              218 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of the V34 strain of cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GP;
                                                                                                                                         Mcmaster JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mcmaster JR,
                                                                                                                                                                                                                                    94US-00367789
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                                                                                                                                                                                                                                                                                                                                                                                                                    virus; strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cucumber
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mosaic
                                                                                                                                            Tricoli DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2; 1
1.7e-06;
hes 0;
                                                                                                                                            Reynolds JF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CMV; V27; V33; V34; A35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reynolds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                            Carney KJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Carney
  to produce which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to produce which are
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ξ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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RESULT 10
ABB80503
ID ABB80503
XX ABB80
XX ABB80
XX ABB80
XX AGrob
  RESULT 11
AAU40661
ID AAU4
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                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based on known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                    This sequence represents a protein sequence from Agrobacterium tumefaciens used within the scope of the invention. The invention relates
                                                                                                                                                                                                                                                                                                                   Sequence 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anti-cucumber mosaic virus tomato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-577926/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sohn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-OCT-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KR99075101-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Agrobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Agrobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABB80503;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABB80503 standard; protein; 218 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Agrobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       resistant
                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                    l Similarity
12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 MDKSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABL58209.
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                                                                                                                                                          KSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MDKSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       to infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                   AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ą,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tumefaciens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tumefaciens;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tumefaciens protein sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98KR-00009096.
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                                                                                                                                                                                                          85.7%; bul
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                          (dd9
                                                                                                                                                                                                                                                                                                                                                                mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 14; DB 2; L
100.0%; Pred. No. 1.7e-06;
... Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Korean.
                                                                                                                                                                                                                                                                                                                                                                virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      anti-cucumber mosaic virus; transgenic
                                                                                                                                                                                                                                          Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hwang YS;
                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                tomato
                                                                                                                                                                                                                    0.00018;
hes 0;
                                                                                                                                                                                                                                                                   DB 3;
                                                                                                                                                                                                                                                              Length 218
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RESULT 12
AAU67189
ID AAU67189
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AAU40661 standard; protein; 70 AA.

standard; protein; 70

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                                                                                                                                                                                                                                                                                                          Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogenic CC polypeptides. The proteins and their associated DNA sequences are used in the treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SAPHO syndrome (synovitis, acne, compared by proteins) and osteomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central cc. P. acnes is also involved in infections of bone, joints and the central cc. P. acnes is associated with acne vulgaris. A method for detecting the gresence or absence of P. acnes in a patient comprises contacting a cc. Sample with a binding agent that binds to the proteins of the invention can determining the amount of bound protein in the sample. The cc. and determining the amount of bound protein in the sample. The collapseptides may be used as antigens in the production of antibodies specific for P. acnes proteins. These antibodies can be used to downregulate expression and activity of P. acnes polypeptides and ctherefore treat P. acnes infections. The antibodies may also be used as can the protein in the sample, by a contract of the c
                                                                                                                     Query Match
Best Local :
                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-APR-2000; 2000US-0199047P.
02-JUN-2000; 2000US-0208841P.
07-JUL-2000; 2000US-0216747P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay;
                                                                                                                                                                                         Sequence 70 AA;
                                                                                                                                                                                                                                                                 this patent did not form part of the printed specification, obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Propionibacterium acnes polypeptides and nucleic acids useful for vaccinating against and diagnosing infections, especially useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-PSDB; AAS59512.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-616774/71.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Skeiky YAW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           dermatological; osteopathic; neuroprotectant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; SEQ ID NO 1856; 1069pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        L'maisonneuve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-APR-2001; 2001WO-US012865.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Propionibacterium acnes immunogenic protein #1557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CORI-) CORIXA CORP
                                                                                           Local Similarity nes 7; Conserv
52
                                           8 SAGRNRR 14
SAGRNRR 58
                                                                                                                                                                                                                                       int/pub/published_pct_sequences
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                                                                                              Conservative
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e J, Zhang Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                vulgaris.
                                                                                           50.0%; Score 7; DB .
100.0%; Pred. No. 8.
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mitcham JL, Wang SS,
, Jen S, Carter D;
                                                                                                                  DB 4;
5. 8.3;
                                                                                              <u>.</u>
                                                                                                                                       Length 70;
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                                                                                              Indels
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                                                                                              Gaps
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RESULT 13
ABM63708
ID ABM63
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                                                                                                                                                                                                                                                                           pustulosis, hypertosis and osteomyelitis), weitis and endopthalmitis.

P. acnes is also involved in infections of bone, joints and the central nervous system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The polypeptides may be used as antigens in the production of antibodies specific for P. acnes proteins. These antibodies can be used to downregulate expression and activity of P. acnes polypeptides and therefore treat P. acnes infections. The antibodies may also be used as diagnostic agents for determining P. acnes protein, by consequence data for this patent did not form part of the printed specification, but was other or the patent at the printed specification, but was only the patent of the printed specification, but was
                                                                                                                                                                 Matches
                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Propionibacterium acnes vaccinating against and treating acne vulgaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptides. The proteins and their associated DNA sequences as the treatment, prevention and diagnosis of medical conditions of the treatment, prevention and diagnosis of medical conditions of the treatment, prevention and diagnosis of medical conditions of medical conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        uveitis; endophthalmitis; bone; joint; central nervous system; Finflammatory lesion; acne vulgaris; enzyme linked immunosorbent dermatological; osteopathic; neuroprotectant.
          ABM63708 standard; protein; 70
                                                                                                                                                                                                                                  Sequence
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02-JUN-2000; 2000US-0208841P
07-JUL-2000; 2000US-0216747P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   L'maisonneuve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Propionibacterium acnes immunogenic protein #28085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-FEB-2002
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                                                                                                                                                                                                                                                               ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                 Local
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DB; AAS59784.
                                                                                             52
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                                                                                                                                                               Similarity 7; Conserv
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                                                                                                                                                                                                                                  70 AA;
                                                                                                                            SAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 28384; 1069pp; English
                                                                                                                                                                 Conservative
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e J, Zhang Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                        50.0%; Sur
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polypeptides and nucleic acids useful for diagnosing infections, especially useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mitcham JL, Wang SS, Jen S, Carter D;
                                                                                                                                                             ; Score 7; DB 4
%; Pred. No. 8.3
0; Mismatches
                                                                                                                                                                               DB 4;
                                                                                                                                                                                             Length 70;
                                                                                                                                                             ..
                                                                                                                                                               Indels
                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           caused by
                                                                                                                                                               Gaps
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cc additionally encompasses expression vectors and host cells comprising a compructed of the invention; antibodies against polypeptides of the invention; fusion proteins comprising a polypeptide of the invention; a comprising the invention; a comprising T cells prepared the development of P. across in a comprision T cells populations or antigen-presenting cells that express the proteins, T cell populations or antigen-presenting cells that express the comprisions. The polymucleotides can also be used as probes or primers for concleic acid hybridisation. The vaccine composition is useful for the stimulation of an immune response against P. across, or for treating across, and the kit is useful for performing a diagnostic assay. The present comprision is useful for the comprision of an immune response composition is useful for the comprision. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from MIPO at ftp. wippo.int/pub/published_pct_sequences
                                                                                                                                               Matches
                                                                                                                                                                                                                   Query Match
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Zhang Y,
Barth B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 encoding a Propionibacterium acnes protein. The invention also polypeptides encoded by the polynucleotides (ABM35624-ABM64536) immunogenic fragments of P. acnes polypeptides. The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to an isolated polynucleotide (ACF64435-ACF64733) encoding a Propionibacterium acnes protein. The invention also relates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-OCT-2002; 2002WO-US032727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; SEQ ID NO 28384; 1481pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Propionibacterium acnes predicted ORF-encoded polypeptide #28384.
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52
                                                                       8
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, Wang S, Jen S, Lod
, Vallieve-Douglass J;
                                                                                                                                       Similarity 7; Conserv
                                                                SAGRNRR 14
                                                                                                                                                                                                                                                                                          70
                                                                                                                                                                                                                                                                                          A
                                                                                                                                               Conservative
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58
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                                                                                                                                                                             100.0%;
                                                                                                                                                                                 50.0%; Score 7; 1
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Lodes MJ,
                                                                                                                                               <u>.</u>
                                                                                                                                               Mismatches
                                                                                                                                                                             DB 6;
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Benson DR,
                                                                                                                                                                                                                   Length 70;
                                                                                                                                               <u>,,</u>
                                                                                                                                               Indels
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Jones R, Ca
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                                                                                                                                           Gaps
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RESULT 14
ABM37180
                                                                CC The invention relates to an isolated polynucleotide (ACF64435-ACF64733) CC encoding a Propionibacterium acnes protein. The invention also relates to CC polypeptides encoded by the polynucleotides (ABM35624-ABM64356) and to CC immunogenic fragments of P. acnes polypeptides. The invention CC additionally encompasses expression vectors and host cells comprising a CC polynucleotide of the invention; antibodies against polypeptides of the cinvention; antibodies against polypeptides of the composition; a comprising a polypeptide of the invention; a comprising a polypeptide of the invention; a cc polynucleotides, antibodies, fusion proteins, T cells prepared CC via this method; a vaccine composition (comprising P. acnes polypeptides, cr polynucleotides, antibodies, fusion proteins, T cell populations, or CC antigen-presenting cells that express the polypeptide); a method and kit Cf or detecting or determining the presence or absence of P. acnes in a cc patient; and a method for inhibiting the development of P. acnes in a cc patient; and a method for inhibiting the development of P. acnes in a cc patient; and a method for inhibiting the development of P. acnes in a cc patient; The P. acnes polypeptides, polynucleotides, antibodies, fusion proteins, T cell populations or antigen-presenting cells that express the volgaris, or for stimulating an immune response specific for a P. acnes (C proteins. The polynucleotides can also be used as probes or primers for cc mucleic acid hybridisation. The vaccine composition is useful for the stimulation of an immune response against P. acnes, or for treating acne, and the kit is useful for performing a diagnostic assay. The present cc invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from with published pct_sequences
                      Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mitcham JL, Skeiky YAW, Pers.
Zhang Y, Wang S, Jen S, Lod
Barth B, Vallieve-Douglass J;
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                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Acne vulgaris; antiseborrhoeic; dermatological; immunostimulant; immune response; vaccine.
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Search completed: December Job time : 74 secs

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AUTHORS
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Capsicum annuum
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots;
spermatophyta; Magnollophyta; eudicotyledons; core eudicots;
latinide; Solanales; Solanaceae; Capsicum.

1 (bases 1 to 127)
Lee,S., Kim,S.-Y., Chung,Y.-H., Shin,H.-J., Goh,S.-H., Pai,H.-S.,
Hur,C.-G. and Choi,D.
Generation of Expressed Sequence Tags from Hot Pepper (Capsicum
annuum L.) and Sequence Analysis in Relation to Hypersensitive
Response Against Pathogen
Unpublished (2001)
                          5 CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGALG 49
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                                                                                                                                                                                                                                                                      Fax: 82-42-860-4309
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                                                                         Similarity
              CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                                                                                                                                                                                                                 quality sequence stop: 127
Location/Qualifiers
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/cultivar="Hang Keun"
/db_xref="taxxn:4072"
                                                                       84.9%;
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mRNA linear EST 11-SEP-2002 cDNA, mRNA sequence.

Genome Research Center and National Center for Genome Information Korea Research Institute of Bioscience and Biotechnology P.O. Box 115, Yusong, Taejeon, 305-600, Republic of Korea

Mismatches DB 4; I . 6.9e-15; 0 Length 127; Indels 0

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thermostabilization and thermoactivation of thermolabile enzymes trehalose and its application for the synthesis of full length cDNA. Proc. Natl. Acad. Sci. U.S.A. 95 (2), 520-524 (1998) Itoh, M., Kitsunai, T., Akiyama, J., Shibata, K., Izawa, M., Kawai, J. Tomaru, Y., Carninci, P., Shibata, Y., Ozawa, Y., Muramatsu, M., Okazaki, Y. and Hayashizaki, Y. Automated filtration-based high-throughput plasmid preparation system. Genome Res. 9 (5), 463-470 (1999)

Carninci, P. and Hayashizaki, Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan
Tel: 81-45-503-9222
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BB440162 RIKEN full-length enriched, 9 days embryo Mus musculus
CDNA clone D030020I06 3' similar to L19932 Mouse (beta ig-h3) mRNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: genome-res@gsc.riken.jp, URL:http://genome.gsc.riken.jp/
Carninci,P., Nishiyama,Y., Westover,A., Itoh,M., Nagaoka,S.,
Sasaki,N., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.
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Unpublished (2000)
Contact: Yoshihide Hayashizaki
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Hayashizaki,Y
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High-efficiency full-length cDNA cloning. Methods Enzymol. 303,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             visit our web site (http://genome.rtc.riken.go.jp)
                                                                                                                                                                                /clone lib="RIKEN full-length enriched, 9 days embryo"
/notee"Site 1: Sall; Site 2: BamHI; cDNA library was
prepared and sequenced in Mouse Genome Expolopedia
Project of Genome Exploration Research Group in Riken
Genomic Sciences Center and Genome Science Laboratory in
RIKEN. Division of Experimental Animal Research in Riken
prepared by using trehalose thermo-activated reverse transcriptase and subsequently enriched for full-length by
                                                                          contributed to prepare mouse tissues. 1st strand cDNA was primed with a primer [5^{\prime}]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mol_type="mRNA"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                       dev_stage="9 days embryo"
lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Mus musculus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        _type="mRNA"
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REFERENCE
AUTHORS
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BB006007
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Konno, H., Alzawa, K., Akahira, S., Akiyama, J., Arakawa, T., Konno, H., Alzawa, K., Fukuda, S., Fukunishi, Y., Hara, A., Hayatsu, N., Hirozame, T., Hori, F., Ishii, Y., Ishikawa, J., Ishikawa, T., Itoh, M., Izawa, M., Kadota, K., Kagawa, I., Kai, C., Kawai, J., Kikuchi, N., Kiyosawa, H., Kojima, Y., Kondo, S., Koya, S., Kurihara, C., Kiyosawa, H., Kojima, Y., Kondo, S., Koya, S., Kurihara, M., Oda, H., Kojasaka, M., Matsuyama, T., Miki, R., Mizuno, Y., Nakamura, M., Oda, H., Okazaki, Y., Ono, T., Owa, C., Saito, H., Sakai, C., Sato, K., Shibata, X., Shibata, Y., Saito, H., Sakai, C., Sato, K., Shibata, X., Shibata, Y., Suzuki, H., Suzuki, H., Tagawa, A., Shiraki, T., Sogabe, Y., Sugahara, Y., Suzuki, H., Suzuki, H., Tagawa, A., Shiraki, A., Watanabe, S., Yamamura, T., Yamanaka, I., Yano, R., Watahiki, A., Yokota, T., Yoshida, K., Yoshiki, A., Yoshino, M., Muramatsu, M. and Hayashizaki, Y.
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                                                                                                                                                                                  Email: genome-res@gsc.riken.jp, URL:http://genome.gsc.riken.jp/Carninci,P., Nishiyama,Y., Westover,A., Itoh,M., Nagaoka,S., Sasaki,N., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.
Thermostabilization and thermoactivation of thermolabile enzymes by trehalose and its application for the synthesis of full length CDNA. Proc. Natl. Acad. Sci. U.S.A. 95 (2), 520-524 (1998)
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Automated filtration-based high-throughput plasmid preparation system. Genome Res. 9 (5), 463-470 (1999)
Carninci,P. and Hayashizaki,Y.
High-efficiency full-length CDNA cloning. Methods Enzymol. 303,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RIKEN Mouse ESTs (Konno,H., et al.)
Unpublished (2000)
Contact: Yoshkhide Hayashizaki
Laboratory for Genome Exploration Research Group, RIKEN Genomic
Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 81-45-503-9222
Fax: 81-45-503-9216
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                             Please visit our web site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACAAATCTGAATCAACCAGT
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                                                                 Cocation/Qualifiers
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Pred. No.
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                                                                                                                                (http://genome.rtc.riken.go.jp) for
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                                                                                                                                                                                                                                                                                                  Konno, H., Aizawa, K., Akahira, S., Akiyama, J., Arakawa, T., Carninci, P., Endo, T., Fukuda, S., Fukunishi, Y., Hara, A., Hayatsu, N., Hirozane, T., Hori, F., Ishii, Y., Ishikawa, J., Ishikawa, T., Itoh, M., Izawa, M., Kadota, K., Kagawa, I., Kai, C., Kawai, J., Kikuchi, N., Izawa, M., Kadota, K., Kagawa, I., Kai, C., Kawai, J., Kikuchi, N., Kiyosawa, H., Kojima, Y., Kondo, S., Koya, S., Kurihara, C., Kiyosawa, H., Kojima, Y., Kondo, S., Koya, S., Kurihara, C., Kiyosawa, A., Kurihara, C., Kiyosawa, A., Kurihara, C., Ono, T., Owa, C., Saito, H., Sakai, C., Sato, K., Okazaki, Y., Ono, T., Owa, C., Saito, H., Sakai, C., Sato, K., Shibata, Y., Shibata, Y., Shigemoto, Y., Shinagawa, A., Shiraki, T., Shibata, Y., Shibata, Y., Shigemoto, Y., Shinagawa, A., Shiraki, A., Sogabe, Y., Sugahara, Y., Suzuki, H., Tagawa, A., Yasunishi, A., Watahiki, A., Watahiki, A., Watahiki, A., Watahiki, A., Yoshida, K., Yoshiki, A., Yoshino, M., Muramatsu, M. and Hayashiyaki Y., Soshida, K., Yoshiki, A., Yoshino, M., Muramatsu, M. and Hayashiyaki Y.
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BB494132 RIKEN full-length enriched, 13 days embryo stomach Mus
musculus cDNA clone D530037K16 3' similar to L19932 Mouse (beta
ig-h3) mRNA, mRNA sequence.
BB494132
                                                                                                                                                                                                                      Hayashizaki,Y.

RIKEN Mouse ESTs (Konno,H., et al.)
Unpublished (2000)
                             Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Jaj
Tel: 81-45-503-9222
Fax: 81-45-503-9216
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                    Contact: Yoshihide Hayashizaki
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genome-res@gsc.riken.jp,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        contributed to prepare mouse tissues. 1st strand cDNA was primed with a primer [5' GAGAGAGAAGGATCCAAGAGCTCTTTTTTTTTTTTTTTVN 3'], cDNA was
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/lab_host="DH10B"
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  URL:http://genome.gsc.riken.jp/
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428 bp
HV05N02u HV Hordeum vulgare subsp.
3-PRIME, mRNA sequence.
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Automated filtration-based high-throughput plasmid preparation system. Genome Res. 9 (5), 463-470 (1999)
Carninci,P. and Hayashizaki,Y.
High-efficiency full-length cDNA cloning. Methods Enzymol. 303,
                                                                                                                                                                                Hordeum vulgare subsp. vulgare
Hordeum vulgare subsp. vulgare
Eukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                 Zhang, H., Potokina, E., Michalek, W., Graner, A.
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Molecular Markers Group, Department Genbank
Institute of Plant Genetics and Crop Plant Research (IPK)
                                                       Barley ESTs from germinating seeds Unpublished (2002)
                                                                                                                                                                                                                                                                                        CA016832.1 GI:24294176
                                                                                                                                                                                                                                                                                                              CA016832
                                        Contact: Stein Nils
                                                                                                                                                                   Pooideae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACAAATCTGAATCAACCA 27
                                                                                                                                          (bases 1 to 428)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Science Enter and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      contributed to prepare mouse tissues. 1st strand cDNA was
                                                                                                                                                             Triticeae; Hordeum.
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/db_xref="taxon:10090"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             clone_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                dev_stage="13 days embryo"
|lab_host="DH10B"
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18;
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                                                                                                                         Weschke, W.,
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Hordeum vulgare subsp. vulgare
Eukaryota, Viridiplantae, Streptophyta, Embryophyta; Tracheophyta;
Spermatophyta, Magnollophyta, Lillopsida, Poales, Poaceae;
Pooideae, Triticeae, Hordeum.
                                                                                  Email: stein@ipk-gatersleben.de
Insert Length: 500 Std Error: 0.00
Plate: 5 row: N column: 2
                                                                                                                                                                                Molecular Markers Group, Department Genbank
Institute of Plant Genetics and Crop Plant Research (IPK)
Corrensstr. 3, 06466, Gatersleben, Germany
Tel: 039482-5522
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HV05N02r HV Hordeum vulgare subsp.
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Tel: 039482-5522
Fax: 039482-5595
                                                                 Seq primer: M13rev
                                                                                                                                                                                                                                                                           Contact: Stein Nils
                                                                                                                                                                                                                                                                                              Barley ESTs from germinating seeds 
Unpublished (2002)
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Zhang, H., Potokina, E., Michalek, W.,
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18; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Vector: pBluescript SK+; Site_1: EcoRI (5'-end of cDNA); Site_2: KhoI (3'-end of cDNA); Roots were grown for two days on filter paper at room temperature. Due to a cloning artefact caused by the kit, in most cases the EcoRI site is NOT present, as well as the EcoRIadapter used for cloning. To excise the insert, restriction sites upstream EcoRI should be used (e.g. BamHI, SalI, PstI). NOTE: Also due to the cloning system used Blue/white selection for recombinats is not 100% reliable.Average insert size is 1 kb"
                         1. .500
                                          Location/Qualifiers
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/dev_stage="germinating seeds (48-96 h)"
/lab_host="XIIIO-Gold"
organism="Hordeum vulgare subsp. vulgare"
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'clone="HV05N02"
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'db_xref="GABI:349120"
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cultivar="barke"
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cDNA clone HV05N02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Weisshaar B
ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BQ595418 597 bp mRNA linear EST 06-DEC-2002 E012691-024-022-C22-SP6 MPIZ-ADIS-024-developing root Beta vulgaris CDNA clone 024-022-C22 5-PRIME, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: weisshaa@mpiz-koeln.mpg.de
Insert Length: 597 Std Error: 0.00
Plate: 22 row: C column: 22
Seq primer: SP6; CATACGATTTAGGTGACACTATAG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes plant J. 32 (5), 845-857 (2002)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M.,
Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Beta vulgaris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BQ595418.1 GI:26125001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Beta vulgaris
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (bases 1 to 597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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/dev stage="germinating seeds (48-96 h)"
/lab_host="XLIO-Gold"
/lab_host="XLIO-Gold"
/clone_lib="HY"
/clone_lib="H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /cultivar="barke"
/sub_species="vulgare"
/db_xref="GABI:267679"
/db_xref="GABO:112509"
                                     /tissue_type="developing root"
/lab_host="EMDH108"
                                                                                                                                                                                                                                                   line)"
                                                                                                                                                                                                                                                                                     /mol_type="mRNA"
/cultivar="KWS2320 (double haploid, monogerm breeding
clone_lib="MPIZ-ADIS-024-developing root"
                                                                                                                     /db_xref="taxon:161934"
/clone="024-022-C22"
                                                                                                                                                                                                                                                                                                                                                                     organism="Beta vulgaris"
                                                                                                                                                                                                                                                                                                                                                                                                                                                ocation/Qualifiers
                                                                                                                                                                                                     db_xref="GABI:191436"
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100.0%; Pred. No. 18;
tive 0; Mismatches
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/notee"Vector: pCMVSPORT6; Site 1: SalI; Site 2: NotI; CDNA library from sugar beet, llbrary provided by KMS Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact: b.schulz@kws.de; cloning sites SalI-NotI, primer sites and

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REFERENCE
AUTHORS
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KEYWORDS
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Best Local Similarity
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Wen,T.J., Qiu,F., Guo,L., Ashlock,D.A and Schnable,P.S.
Expressed Sequence Tags from B73 Maize: various stages and tissues including seedlings treated with a variety of hormones Unpublished (2001)
Contact: Patrick S. Schnable
Schnable Laboratory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (<a href="http://depts.washington.edu/ventures/collabtr/direct/index.htm>#b rt). Overall sequence quality assessment and vector trimming were conducted using the Lucy software (<a href="http://www.tigr.org/softlab/">http://www.tigr.org/softlab/</a>). Lucy parameters were set to ensure an overall trimmed quality of 97.5% or better without any vector fragments in the chosen high-quality region of each sequence. Low-quality bases between the poly-T and the high-quality region were replaced with N's to serve
                                                                                                                                                                                                                                                                                                                                                      FORWARD: primer T7-1 (AA TAC GAC TCA CTA TAG)
BACKWARD: primer T3 (ATT AAC CCT CAC TAA AG)
Seq primer: primer T3 (ATT AAC CCT CAC TAA AG)
Location/Qualifiers
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Individual basecall and confidence value were assigned using
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/lab_nose="ISUNS-RN"
/clone_lib="ISUNS-RN"
/clone_rib="ISUNS-RN"
/note="Vector: pT7T3PAC; Site_1: EcoRI; Site_2: NotI;
/note="Vector: pT7T3PAC; Site_1: EcoRI; Site_2: NotI;
Tissues: Germinated seed and seedlings (1, 2, 8, 11 DAG),
Tissues: Germinated seed (17, 21, 38, 69, 77 DAG), Kernels
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                                                                                                                                  /clone="MEST183-H08"
/tissue_type="mixed"
/lab_host="DH108"
                                                                                                                                                                                                                                                                     /organism="Zea mays"
/mol_type="mRNA"
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                                                                                                                                                                                                                                                    cultivar="B73"
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Pred. No.
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VERSION KEYWORDS

ACCESSION

JOURNAL COMMENT

FORWARD: primer T7-1 (AA TAC GAC TCA CTA TAG)
BACKWARD: primer T3 (ATT AAC CCT CAC TAA AG)

PCR PRimers

REFERENCE

AUTHORS TITLE

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SOURCE
ORGANISM
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(<a href="http://depts.washington.edu/ventures/collabtr/direct/index.htm=#brt)">http://depts.washington.edu/ventures/collabtr/direct/index.htm=#brt)</a>. Overall sequence quality assessment and vector trimming were conducted using the Lucy software (<a href="http://www.tigr.org/softlab/">http://www.tigr.org/softlab/</a>). Lucy parameters were set to ensure a noverall trimmed quality of 97.5% or better without any vector fragments in the chosen high-quality region of each sequence. Low-quality bases between the poly-T and the high-quality region were replaced with N's to serve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wen,T.J., Qiu,F., Guo,L., Ashlock,D.A and Schnable,P.S. Expressed Sequence Tags from B73 Maize: various stages including seedlings treated with a variety of hormones Unpublished (2001)
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EST.
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MEST279-G08.T3 ISUM5-RN Zea mays cDNA clone MEST279-G08 3', mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Schnable Laboratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Patrick S. Schnable
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515-294-0975
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18;
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RESULT 9

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ORIGIN

Matches

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REFERENCE
AUTHORS
TITLE
JOURNAL
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AG223668
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Best Local Similarity
                                                         JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18;
                                                                                                                                     Sato, S., Nakamura, Y. and Tabata, S. Lotus japonicus BAC End sequences Published Only in Database (2002) 2 (bases 1 to 627)
                                                                                                                                                                                                                                                                                                    Lotus corniculatus var. japonicus (Lotus japonicus)
Lotus corniculatus var. japonicus
Lotus corniculatus var. japonicus
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots;
rosids; eurosids I; Fabales; Fabaceae; Papillonoideae; Loteae;
                          Submitted (20-NOV-2002) Shusei Sato, Kazusa DNA Research Institute, The First Laboratory for Plant Gene Research; 2-6-7
                                                                                   Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AG223668.1 GI:26532397
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lotus corniculatus var. survey sequence.
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Kazusa-kamatari, Kisarazu, Chiba 292-0818, Japan
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Location/Qualifiers
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llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tissues: Germinated seed and Seedlings (1, 2, 8, 11 DAG), Mixed mature tissures (17, 21, 38, 69, 77 DAG), Kernels (3, 5, 10, 15, 20, 25, 30, DAP), Adventious roots (65 DAG), Tassel (3-39 cm, 53 and 56 DAG), Immature ear (0.2-3.0 cm, 53, 56, 59 DAG), Husk (73 DAG), Silk, unpollinated first ear, ear shank, etiolated seedlings, cycloheximide-treated callus, Anaerobic treated seedlings, NAA (a-Naphthalene acetic acid)-treated seedlings, Kinetin-treated seedlings, KOPC (1-aminocyclopropane-1-carboxylix acid)-treated seedlings, Brassinolide-treated seedlings, ABA (Abscisic acid)-treated seedlings, JA (Jasmonic acid)-treated seedlings, JA (Jasmonic acid)-treated seedlings, Seedlings, JA (Jasmonic acid)-treated seedlings, JA (Jasmonic acid)-treated seedlings of cDNA was prepared from oligo-dT selected mRNA by priming with a Notic oligo-dT primer (5)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      resulting DNA:RNA hybrid was treated with RNase H and used as a template for DNA PolI-catalyzed second strand synthesis. After the addition of EcoRI adaptors, the ds-cDNAs were digested with NotI and size-selected. The resulting molecules were directionally cloned into the EcoRI and NotI sites of the pT773PAC vector. The library then went through one round of normalization to CoT value of 5 based on the methods of Marcelo Bento Soares (Genome Research 6: 791-806, 1996)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="ISUM5-kw"
/note="Vector: pT7T3PAC; Site_1: EcoRI; Site_2: NotI;
/note="Vector: pT7T3PAC; Site_1: EcoRI; Site_2: NotI;
//isues: Germinated seed and seedlings (1, 2, 8, 11 DAG),
Tissues: Germinated seed and seedlings (1, 2, 8, 11 DAG),
//isues-assumes-seed (17, 21, 38, 69, 77 DAG)
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Pred. No.
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Collect. ..., Plant Biology Division
The Samuel Roberts Noble Foundation
The Samuel Roberts Noble Foundation
OK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Medicago truncatula
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NF014E05FL1F1038 Developing flower Medicago truncatula cDNA clone
NF014E05FL 5', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: gdmay@noble.org
Insert Length: 629 Std Error: 0.00
Plate: 014 row: E column: 05
Seq primer: TCACACAGGAAACAGCTATGAC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 580 224 6650
Fax: 580 224 6692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Torres-Jerez,I., Scott,A.D., Harris,A.R., Gonzales,R.A., Bell,C., Flores,H.R., Inman,J.T., Weller,J.W. and May,G.D. Expressed Sequence Tags from the Samuel Roberts Noble Foundation Medicago truncatula flower library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Trifolieae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: May GD
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/dev stage="Developing filly-opened flowers and flowers in early transition into pods."
/clone_ilb="Developing flower"
/clone_ilb="Developing flower"
/clone_ilb="Developing flower"
/clone_ilb="Developing flower"
/note="Vector: Lambda Zap; cDNA was prepared from polyA+enriched, pooled samples of equivalent amounts of total RNA from very young, developing, fully-opened flowers and flowers transitioning into pods. The cDNA was directionally ligated into the Uni-Zap XR vector (Stratagene) and packaged using the Glapapack III Gold packaging extracts. Phagemids containing cDNA inserts were in vivo excised from the recombinant Uni-Zap XR vector using ExAssist helper phage and the E. coli strain
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/clone_lib="genomic BAC library"
/note="VECTOR:pBeloBAC11~synonym: Lotus japonicus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="genomic DNA"
/strain="Miyakojima MG-20"
/variety="japonicus"
/db_xref="taxon:34305"
                                                                                                                                                                                                                                                                                                                                  /clone="NF014E05FL"
/tissue_type="Developing_flowers"
                                                                                                                                                                                                                                                                                                                                                                                       /mol_type="mRNA"
/db_xref="taxon:3880"
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                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Medicago truncatula"
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USA

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RESULT 13
CK185060
LOCUS
DEFINITION
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AUTHORS
TITLE
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BX752014
LOCUS
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Best Local Similarity
Matches 18; Conserv
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                                                                                                                          332
CK185060 688 bp mRNA linear EST 01 EST774375 BEA Boophilus microplus cDNA clone BEAC260, mRNA
                                                                                                                                                 3 TCCATGGACAAATCTGAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence is from a Xenopus Gene Collection (XGC) library constructed by Aaron M. Zorn.

CDNA was oligo dI primed from Sug of poly A+ RNA from stages 10-13 gastrulae. EcoRI-NotI cut cDNA was then ligated into pCS107 with EcoRI at the 5' end and NotI at the 3' end.

Vector: pCS107; Site 1: EcoRI; Site 2: NotI
Host: Escherichia coli X1- blue.

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 688)
Croning, M.D.R., Ashurst, J.L., Taylor, R., Zorn, A.M. and Rogers, J. Sanger Xenopus tropicalis EST project 2001 (11_2003)
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BX752014 XGC-gastrula Xenopus tropicalis cDNA clone TGas076c13 3',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GATCCATGGACAAATCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: trop@sanger.ac.uk
Sanger Xenopus tropicalis EST project 2001
TROPICALIS_SEQUENCE_ID: TGas076cl3.qlkT7
Sequencing_primer: T7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Xenopus tropicalis (western clawed frog) Xenopus tropicalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sanger Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Croning MDR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hinxton, Cambridgeshire, CB10 1SA, UK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kenopodinae; Xenopus; Silurana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GATCCATGGACAAATCTG 91
                                                                                                                          TCCATGGACAAATCTGAA 349
                                                                                                                                                                                                 34.0%; Score 18; DB llarity 100.0%; Pred. No. 18 Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                     /dev_stage="gastrula (stages 10.5-12 mixed)"
/lab_host="Escherichia coli XLI-blue"
/clone lib="XGC-gastrula"
/note="Vector: pCS107; Site_1: EcoRI; Site_2: NotI; cDNA
was oligo dT primed from Sug of poly A+ RNA from stages
10-13 gastrulae. EcoRI-NotI cut cDNA was then ligated
into pCS107 with EcoRI at the 5' end and NotI at the 3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    XL1-Blue MRF' (Stratagene). Excised plasmids were plated using SOLR cells. \ensuremath{^{\rm T}}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Xenopus tropicalis"
/mol_type="mRNA"
/db_xref="taxon:8364"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  clone="TGas076c13"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           34.0%; 50
100.0%; Pr
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Pred. No.
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                     EST 01-JUL-2004
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AUTHORS
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AUTHORS
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CB634510/c
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SOURCE
ORGANISM
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                                                                                                                                               COMMENT
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                                                                                                                                                                                                                                                                             Oryza sativa (indica cultivar-group)
Oryza sativa (indica cultivar-group)
Oryza sativa (indica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermattophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                              CB634510
OSIIEb14B06.f OSIIEb Oryza sativa (indica cultivar-group) cDNA clone OSIIEb14B06.5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18;
                                                                                                                                                           Jantasuriyarat, C., Lu, G., Gowda, M., Hatfield, J., Zhou, B., Mazur, E., Kudrna, D., Dean, R., Soderlund, C., Wing, R. and Wang, G. Large-scale identification of ESTs involved in the interaction between rice and Magnaporthe grisea
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                         clone OSIIEb14B06 5', ml
CB634510
CB634510.1 GI:29629501
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Other_ESTs: EST774376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence.
CK185060
CK185060.1 GI:49565594
Tel: 520 626 3967 Fax: 520 621 9288 Email: http://genc
                                                                           Arizona Genomics Institute
University of Arizona
Biological Sciences West, '
                                                                                                                                                                                                                                                                                                                                                                                         EST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: nene@tigr.org
Seq primer: M13 forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The Institute for Genomic Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Vishvanath Nene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Boophilus microplus
                                                               85721-0088,
                                                                                                                                             Contact:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fax: 301-838-0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9712 Medical Center Drive, Rockville,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Parasite Genomics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Boophilus microplus (southern cattle tick)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                (bases 1 to 701)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
  http://genome.arizona.edu
                                                                                                                                           Rod Wing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="BEA"
/note="Organ: Whole ticks and dissected organs; Vector:
/note="Organ: Whole ticks and dissected organs; Vector:
pExpress | 1 A normalized cDNA library was custom built by
Express Genomics. Oligo-dT primed cDNA was directionally
cloned into NotI-EcoRV site of pExpress 1. Universal M13
primers were used to generate 5' and 3' EST data."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Boophilus microplus"
/mol_type="mRNA"
/db_xref="taxon:6941"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /dev_stage="Adult and larvae"
/lab_host="E. coli strain DH10B-TonA"
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                                                                                West, 448A,
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                                                                                  P.O.
                                                                                  Box 210088, Tucson,
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EST 08-APR-2003

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SOURCE
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CC501966/c
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AUTHORS
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VERSION
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         586
                                                                                                                                        Email: rholt@bcgsc.ca
Email: rholt@bcgsc.ca
Clones are derived from the bovine BAC library CHORI-240
(http://www.chori.org/bacpac/bovine240.htm). For BAC library
availability, please contact Pieter de Jong (pdejong@mail.cho.org).
Clones may be purchased from BACPAC Resources
(http://www.chori.org/bacpac/ordering_information.htm). This work
was undertaken as part of the International Bovine BAC Mapping
Consortium (IBBMC) by CSIRO Livestock Industries, Australia and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC501966 713 b
CH240 340H21.T7 CHORI-240 Bos
genomic survey sequence.
CC501966
CC501966 1 GI;31820259
                                                                                                                                                                                                                                                                                                                                                                      Sequencing
The British Columbia Cancer Agency Genome Science Centre
600 W. 10th Ave, Vancouver, British Columbia, Canada V5Z
Tel: 604-877-6085
Fax: 604-877-6276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Holt, R., Stott, J., Yang, G., Barber, S., Smailus, D., Prabhu, A.-L., Tsai, M., Cloutler, A., Lee, D., Girn, N., Olson, T., Mayo, M., Butterfield, Y., Kirkpatrick, R., Liu, J., Guin, R., Chan, A., Chiu, R., Mathewson, C., Wye, N., Masson, A., Brown-John, M., Jones, S., Schein, J., Marra, M., de Jong, P., McWilliam, S., Barris, W., Dalrymple, B.P. and Tellam, R. Bovine BAC End Sequences from Library CHORI-240, PLATES 294 to 398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GSS.
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FORWARD: gta aaa cga cgg cca gtg
BACKWARD: gga aac agc tat gac cat g
Plate: 14 row: B column: 06
Seq primer: gta aaa cga cgg cca gtg.
                                                                                  British Columbia Genome Sciences Centre, Plate: 340 row: H column: 21 Seq primer: T7 Class: BAC ends.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2003)
Other_GSSs: CH240_340H21.TARBAC13P2
Contact: Rob Holt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bos taurus (cow)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (bases 1 to 713)
1. ./13
/organism="Bos taurus"
                                                          Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone lib="OSIIEb"
/note="Vector: pBluescript II KS +; Site_1: EcoRI; Site_2:
KhoI; 24 hrs after innoculation with Rice Blast (PO6-6-3)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tissue_type="Leaf"
/dev_stage="3 week"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              organism="Oryza sativa (indica cultivar-group)"
|mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       db xref="taxon:39946"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lone="OSIIEb14B06"
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100.0%; Pred. No. 18;
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taurus genomic clone CH240_340H21,
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Search completed: December 5, 2004, 07:59:11 Job time : 1487 secs
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                                                                                                                                                                                                                                                                                                                                          /mol type="genomic DNA"
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/db xref="taxon:9913"
/clone="CH240_340H21"
/sex="Male"
                                                                                                                                                                                                                                             /cell_type="Blood"
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/clone lib="CHORI-240"
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/note="Vector: pTARBAC1.3; Site 1: MboI; Site 2: MboI;
Hereford bull Li Domino 99375; CHORI-240 Bovine BAC
library (Male) produced by Pieter de Jong"
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100.0%; Pred. No.
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GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd

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ALIGNMENTS	A41823	E01821	MCVCPA	CMO131623	CMO131624	AR076864	A67212	CMVCOATP	CMU22821	CUMCVRN4	BD000279	AR430190	AR364711	AR138343	AR021487	A39812	A32133	BD000278	AR364710	AR138342	AR021486	A32131	MCVCPB	CMU32858	CMU32859	I49963
	A41823 Sequence 4	E01821 cDNA encodi	M22710 Cucumber mo	AJ131623 Cucumber	AJ131624 Cucumber	AR076864 Sequence	A67212 Sequence 1	X77855 Cucumber Mo		X16386 Cucumber mo	BD000279 Transgeni	AR430190 Sequence	AR364711 Sequence	AR138343 Sequence	AR021487 Sequence	A39812 Sequence 6	A32133 CMV capsid	BD000278 Transgeni	AR364710 Sequence	AR138342 Sequence	AR021486 Sequence	A32131 CMV capsid	L36525 Cucumber mo	U32858 Cucumber mo	U32859 Cucumber mo	149963 Sequence 13

RESULŤ 1 BD261798 LOCUS DEFINITION

BD261798 53 bp DNA linear PAT 17-UTL-2003 Enhancement in protein production by higher plants using ubiquitin or cucumber mosaic virus coating protein peptide.

BD261798 FREEDERS OF THE PROTECT OF THE P PAT 17-JUL-2003

ACCESSION VERSION KEYWORDS BD261798.1 GI:33071566 JP 2002532098-A/2

SOURCE ORGANISM Cucumber mosaic virus (cucumber mosaic cucumovirus)
Cucumber mosaic virus

REFERENCE AUTHORS TITLE Cucumovirus. Viruses; ssRNA positive-strand viruses, no DNA stage; Bromoviridae;

1 (bases 1 to 53)
1 (bases 1 to 53)
1 (Pang,R.X., Wu,J.L. and Chen,X.Y.
Frang,R.X., Wu,J.L. and Chen,X.Y.
Enhancement in protein production by higher plants using ubiquitin for cocumber mosaic virus coating protein peptide patent: JP 2002532098-A 2 02-OCT-2002;
INSTITUTE OF MOLECULAR AGROBIOLOGY

PP PR OS

COMMENT

JOURNAL

Cucumber mosaic virus
JP 2002532098-A/2
O2-OCT-2002
11-DEC-1998 JP 2000588378
RONG XIANG FANG, JUNG LIN WU, XIAO YING CHEN
C12N15/09, A01H5/00, C07K14/415, C07K19/00, C12N5/10, C12N15/00, PC

C12N5/00

Enhancement in protein production by higher plants using ubiquitin or

protein peptide FH

Ξ cucumber mosaic virus coating Location/Qualifiers

source (6). (47). Location/Qualifiers

FEATURES

/organism="Cucumber mosaic virus" /mol\_type="genomic DNA" /db\_xref="taxon:12305"

Query Match Best Local S Similarity 53; Conserv 100.0%; silarity 100.0%; F Conservative 0; Score 53; DB 6; 1 Pred. No. 2.2e-20; ; Mismatches 0; Length 53;

ORIGIN

1 GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACGAGCT 53

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GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACGAGCT
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TITLE JOURNAL FEATURES KEYWORDS SOURCE ORGANISM DEFINITION ACCESSION VERSION KEYWORDS REFERENCE AUTHORS REFERENCE AUTHORS RESULT 2 AR111176 LOCUS SOURCE ORGANISM RESULT 4 AR111975 DEFINITION ACCESSION RESULT 3 AR111981 밁 S DEFINITION ACCESSION ORIGIN 밁 á ORIGIN FEATURES SOURCE VERSION POCUS VERSION **CEYWORDS** Snoo Query Match Best Local S Matches 46 Query Match Best Local S Matches 46 TITLE JOURNAL ORGANISM source Bource AR111975 Sequence 3 : AR111975 AR111975.1 Carney, K.J.

Plants resistant to C strains of cucumber mosaic virus

Patent: US 6127601-A 14 03-CCT-2000;

Location/Qualifiers

1. 772

1. 772 4 46; AR111981 Sequence 14 from patent AR111981 1 (bases 1 to 772) Boeshore, M.L., McMaster, R.J., Tricoli, D.M., 46; Carney, K.J.

Plants resistant to C strains of cucumber mosaic virus patent: US 6127601-A 5 03-OCT-2000; 1 (bases 1 to 771)
Boeshore, M.L., McMaster, R.J., Tricoli, D.M., Unknown AR111976.1 Sequence 5 from patent US AR111976 Unknown. Unknown. Unknown. Unknown. AR111976 Unclassified. AR111981.1 Similarity n 86.8%; Score 46; DB Similarity 100.0%; Pred. No. 3.4 46; Conservative 0; Mismatches Unclassified. CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49 CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49 CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 46 86.8%; Score 46; DB ilarity 100.0%; Pred. No. 3.6 Conservative 0; Mismatches /organism="unknown" /mol\_type="unassigned DNA" organism="unknown" Location/Qualifiers from patent US GI:12828829 GI:12828824 GI:12828823 \_type="unassigned DNA" 772 bp US 6127601. 792 bp S 6127601. 771 bp 6127601. DB 6; L., 3.6e-16; 0; DB 6; Leng 3.6e-16; DNA DNA Length 772 Length 771; Indels Indels linear linear linear Reynolds, J.F. Reynolds, J.F. and 46 PAT PAT PAT 0 0 14-FEB-2001 14-FEB-2001 14-FEB-2001 Gaps 0 0

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REFERENCE
AUTHORS
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AR031575
                                                                                                           1 (bases 1 to 130)
Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.
Cell ablation using trans-splicing ribozymes
Patent: US 5849548-A 14 15-DEC-1998;
Location/Qualifiers
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5866394-A 14 02-FEB-1999;
Location/Qualifiers
 L Similarity 45; Conserv
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1 (bases 1 to 792)

1 Boeshor, M.L., McMaster, R.J.,
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84.9%; Score 45; DB 6; Lilarity 100.0%; Pred. No. 1.3e-15; Conservative 0; Mismatches 0;
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/mol_type="unassigned DNA"
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Sequence 11 from patent
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Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M. Cell ablation using trans-splicing ribozymes Patent: US 5641673-A 14 24-UUN-1997;
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Haseloff,J., Brand,A., Perrimon,N. a
Cell ablation using trans-splicing r
Patent: US 6071730-A 14 06-JUN-2000;
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1 (bases 1 to 131)
Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M. Cell ablation using trans-splicing ribozymes
Patent: US 6071730-A 11 06-JUN-2000;
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M. Cell ablation using trans-splicing ribozymes
Patent: US 5849548-A 11 15-DEC-1998;
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M. Cell ablation using trans-splicing ribozymes
Patent: US 5866384-A 11 02-FEB-1999;
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Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5641673-A 11 24-UUN-1997;
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Sequence 11 from patent US 5641673.
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Sequence 9 from patent
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M.
Cell ablattion using trans-splicing ribozymes
Patent: US 5866384-A 9 02-FEB-1999;
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Similarity 100.0%; Pred. No. 1.3e-15;
45; Conservative 0; Mismatches 0;
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AR097441
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Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 6071730-A 9 06-JUN-2000;
Location/Qualifiers
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Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5849548-A 9 15-DEC-1998;
Location/Qualifiers
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Result
No.
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Word size :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OM nucleic - nucleic search, using sw model
                                                    C
                                                                                                                                                                                                                                                         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                    Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                seq length: 0
seq length: 2000000000
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Match
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Gapop 60.0 , Gapext 60.0
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|: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*
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/cgn2_6/ptodata/2/pubpna/US09C_PUBCOMB.seq:*
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/cgn2_6/ptodata/2/pubpna/US10E_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US10E_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US10D_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US10E_PUBCOMB.seq:*
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cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
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960
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            US-10-011-033-5
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US-10-425-115-38826
US-10-425-115-38826
US-10-426-761-6291
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US-10-126-761-6291
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## ALIGNMENTS

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Sequence 5, Appli
Sequence 14, Appli
Sequence 3, Appli
Sequence 1, Appli
Sequence 9, Appli
Sequence 35833, A
Sequence 38826, A
Sequence 4291, Ap
Sequence 94969, A
Sequence 94969, A
Sequence 950, Ap
Sequence 97141, A
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US-10-011-033-5
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Publication No. US20020124286A1
GENERAL INFORMATION:
APPLICANT: Boeshore, Maury L
                                                                                                                             COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-NO. US20020124286A1-2001
CLASSIFICATION: CUnknown>
                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-UN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Carney, Kim u
TITLE OF INVENTION: Plants Resistant to C
Cucumber Mosaic Virus
              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                   STREET: 2 Prudent
CITY: Chicago
STATE: Illinois
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson,
                                                                                                                                                                                                                                                                                                                        60089
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Tricoli, David M
Reynolds, John F
Carney, Kim J
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US-10-011-033-14
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                 TELEFAX: 312-616-5460 INFORMATION FOR SEQ ID NO: 14:
                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/10/011,033
APPLICATION NUMBER: US/20020124286A1-2001
CLASSIFICATION ENAMOND
PRIOR APPLICATION NUMBER: US/08/875,233
FILING DATE: 3-UN-1997
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELEPHONE: 312-616-5400
TELEPHONE: 312-616-5400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                / Match 86.8%; Score 46; DB 13; Local Similarity 100.0%; Pred. No. 1.7e-16 es 46; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 771 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Plants Resistant to C Strains Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
Carney, Kim J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /standard_name= "COAT PROTEIN" SEQUENCE DESCRIPTION: SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
SEQUENCE CHARACTERISTICS:
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LOCATION: 3..660
OTHER INFORMATION: /codon start= 3
/function= "ENCAPSIDATES VĪRUS RNA"
/product= "COAT PROTEIN"
/gene= "CP"
                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.
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hes 0;
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Publication No. US20020124286A1
GENERAL INFORMATION:
APPLICANT: Boeshore, Maury L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                 ANTI-SENSE: NO ORIGINAL SOURCE:
                                                         MOLECULE TYPE: cl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HYPOTHETICAL: NO ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        46;
                                                                                                                   LENGTH: 792 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 2 Pruder
CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                  TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Boeshore, Maury L
                                                                                 CDNA
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ORIGINAL SOURCE:
ORGANISM: Cucumber Mosaic Virus
INDIVIDUAL ISOLATE: A35
                      4 CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 46
                                                                          86.8%; Score 46; DB 13; 1
100.0%; Pred. No. 1.7e-16;
/ative 0; Mismatches 0;
                                                                                                                                                                              14:
                                                                                                                Length 772;
                                                                            Indels
                                                                            0;
                                                                            Gaps
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; SEQUENCE DESCRIPTION: SEQ ID NO: US-10-011-033-14 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5460
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS: COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARBE: PATECRETA; PC-DOS/MS-DOS
CURRENT APPLICATION DATA: Release #1.0, Version #1.30
APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION: <Unknown> McMaster, J. Russell Tricoli, David M Reynolds, John F Carney, Kim J TITLE OF INVENTION: Plants Resistant to C Strains COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk ORGANISM: CUCUMBER MOSAIC VIRUS ADDRESSEE: Rockey, Milnamow & Katz, Ltd. STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700 Cucumber Mosaic Virus

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; NAME/KEY: CDS
; LOCATION: 3..660
; SEQUENCE DESCRIPTION: S
US-10-011-033-1
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; LOCATION: 3..660
; SEQUENCE DESCRIPTION: S
US-10-011-033-3
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US-10-011-033-1
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APPLICATION NUMBER: US/10/011,033

FILING DATE: 13-No. US20020124286A1-2001

CLASSIFICATION: <UNLOwn
PRIOR APPLICATION: <UNLOwn
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/875,233

FILING DATE: 26-JUN-1997

ATTORNEY/AGENT INFORMATION:

NAME: Lisa V. Mueller

TELECOMMUNICATION INFORMATION:

TELEPHONE: 312-616-5460

INFORMATION FOR SEQ ID NO: 1:

SEQUENCE CHARACTERISTICS:
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/10011033 Publication No. US20020124286A1 GENERAL INFORMATION:
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                                                                                                                                                                                                       LENGTH: 772 base pairs
TYPE: nucleic acid
STRANDENMESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Reynolds, John r
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains of
Cucumber Mosaic Virus
                                                                                                                                                ORIGINAL SOURCE:
ORGANISM: Cucumber Mosaic Virus
INDIVIDUAL ISOLATE: V-27
                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           46;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN: v-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     McMaster, J. Russell
Tricoli, David M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Boeshore, Maury L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 Prudential Plaza, 180 N. Stetson, Suite 4700
 81.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               86.8%; Score 46; DB 13; Length 792; 100.0%; Pred. No. 1.7e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO: 3:
                                                                          SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>,,</u>
Score 43; DB 13;
Pred. No. 8.9e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                   Length 772;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
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                                                                                                                                                        US-10-011-033-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9, Application US/10011033 Publication No. US20020124286A1 GENERAL INFORMATION:
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                                                                           Matches
                                                                                               Best
                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION EVENTON: VUNKNOWN>
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-UN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECHONE: 312-616-5460
TELECHONE: 312-616-5460
                                                                                             Local
                                                                                                                                                                                                                                              ORIGINAL SOURCE:
ORGANISM: CUCUMBER MOSAIC VIRUS
STRAIN: STRAIN C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                         SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Plants Resistant to C Strains Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                         ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Boeshore, Maury L
                                                                       ch 69.8%; Score 37; DB 13; Length 960; Similarity 100.0%; Pred. No. 2.6e-11; 37; Conservative 0; Mismatches 0; Indels
                                6 ATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACC 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCG 46
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                                                                                                                                                                                           NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 960 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CCATGGACAATCTGAATCAACCAGTGCTGGTCGTAACCGTCG 43
ATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACC 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: Illinois
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Reynolds, John Carney, Kim J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  McMaster, J. Ru
Tricoli, David
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      J. Russell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ٥,
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                                                                                                                                                                         9:
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                                                                           0;
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RESULT 6 US-10-369-493-35833/c

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APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN TITLE OF INVENTION: DAINS: WITH IMPROVED PROPERTIES FILE REFERENCE: 38-10(52052)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
                                                                                                                                                                     RESULT 8
US-10-156-761-6291/c
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US-10-369-493-35833
APPLICANT: OMURA, SATOSHI
APPLICANT: IKEDA, HARUO
APPLICANT: ISHIKAWA, JUN
APPLICANT: HORIKAWA, HIROS
APPLICANT: SHIBA, TADAYOSH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 38826
LENGTH: 1020
                                                                                                          GENERAL INFORMATION:
                                                                                                                              Sequence 6291, Application US/10156761 Publication No. US20030119018A1
                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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LENGTH: 1674
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: La ROBA, Thomas J. APPLICANT: Kovalic, David K. APPLICANT: Zhou, Yihua APPLICANT: Zhou, Yihua APPLICANT: Cao, Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 47374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local 20;
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                                                                                                                                                                                                                                                            983 TGAATCAACCAGTGCTGG 1000
                                                                                                                                                                                                                                                                                        17 TGAATCAACCAGTGCTGG 34
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                                                                                                                                                                                                                                                                                                                                              18;
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                      IKEDA, HARUO
ISHIKAWA, JUN
HORIKAWA, HIROSHI
  SHIBA, TADAYOSHI
                                                                                                                                                                                                                                                                                                                                              Conservative
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100.0%; F:
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100.0%; Pred. No.
tive 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                               Score 18; DB 18; Length 1020; pred. No. 2.3;
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CURRENT APPLICATION NUMBER: US/10/156,761
CURRENT FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: JP 2001-204089
PRIOR FILING DATE: 2001-05-30
PRIOR APPLICATION NUMBER: JP 2001-272697
PRIOR APPLICATION NUMBER: JP 2001-272697
PRIOR FILING DATE: 2001-08-02
NUMBER OF SEQ ID NOS: 15109
SEQ ID NO 6291
LENGTH: 1116
                                                                                                                                                                                                                 RESULT 10
US-10-101-464A-950/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_56769C.1
US-10-424-599-94969
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Chou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE REFERENCE: 38-2153223)B
TITLE REFERENCE: 38-2153223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 94969
LENGTH: 1996
                  Sequence 950, Application US/10101464A
Publication No. US20030046728A1
GENERAL INFORMATION:
APPLICANT: Strabala, Timothy
APPLICANT: Nieuwenhuizen, Nicolaas
APPLICANT: Higgins, Colleen M.
TITLE OF INVENTION: Compositions Isolated from Plant Cells
TITLE OF INVENTION: and Their Use in the Modification of Plant Cell Signaling
FILE REFERENCE: 11000.1020c2
                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 94969, Application US/10424599 Publication No. US20040031072A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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APPLICANT: HATTORI, MASAHIRA
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-262
CURRENT APPLICATION NUMBER: US/10/101,464A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Streptomyces avermitilis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: CDS
LOCATION: (1).
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                                                                                                                                                                                                                                                                                                         532
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                                                                                                                                                                                                                                                                                                                                               33 GGTCGTAACCGTCGACGA 50
                                                                                                                                                                                                                                                                                                                                                                                          18;
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                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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. 2.3;
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PRIOR APPLICATION NUMBER: 09/704,302
PRIOR FILING DATE: 2000-11-01
PRIOR FILING DATE: 1909-01-12
PRIOR FILING DATE: 1999-01-12
PRIOR APPLICATION NUMBER: 60/162,866
PRIOR FILING DATE: 1999-11-01
PRIOR APPLICATION NUMBER: FCT/US00/00724
PRIOR APPLICATION NUMBER: PCT/US00/00724
PRIOR FILING DATE: 2000-01-11
NUMBER OF SEQ ID NOS: 989
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 950
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; ORGANISM: Eucalyptus grandis
US-10-101-464A-950
US-10-437-963-97141, Application US/10437963; Sequence 97141, Application US/20040123343A1; Publication No. US20040123343A1; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.; APPLICANT: Kovalic, David K.
                                                                                                                                                                                           밁
                                                                                                                                                                                                                                                                                                                                                            ; PEATURE:

, NAME/KEY: misc feature

; LOCATION: (4187715)

; OTHER INFORMATION: a, t, c,

US-10-156-761-1
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US-10-156-761-1
                                                                                                                              RESULT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SAKAKI, YOSHIYUKI
APPLICANT: HATTORI, MASAHIRA
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-262
CURRENT APPLICATION NUMBER: US/10/156,761
CURRENT FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: UP 2001-204089
PRIOR APPLICATION NUMBER: JP 2001-272697
PRIOR APPLICATION NUMBER: JP 2001-272697
PRIOR APPLICATION NUMBER: JP 2001-272697
PRIOR FILING DATE: 2001-08-02
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APPLICANT: IKEDA, HARUO
APPLICANT: ISHIKAWA, JUN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 15109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 9025608
TYPE: DNA
ORGANISM: Streptomyces avermitilis
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HORIKAWA, HIROSHI
SHIBA, TADAYOSHI
                                                                                                                                                                                                                                                                       34.0%; Score 18; DB 15; Length 9025608; ilarity 100.0%; Pred. No. 1.8; Conservative 0; Mismatches 0; Indels 0;
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US-10-027-632-3534
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CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
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APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Wu, Wei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT APPLICATION NUMBER: US/10/437,963
SEQ ID NO 97141
LENGTH: 285
LENGTH: 285
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SEQ ID NO 3534
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Best Local
                                                                            Best Local Similarity Matches 17; Conserv
                                                                                                                        Query Match
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/185,218 PRIOR FILING DATE: 2000-02-24
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LOCATION: (1)..(285)
OTHER INFORMATION: unsure at all n locations
                                                                                                                                                                                        TYPE: DNA
ORGANISM: Human
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474 AAATCTGAATCAACCAG
                                       12 AAATCTGAATCAACCAG 28
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17; Conserv
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100.0%; Pr
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490
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Pred. No. 9.1;
0; Mismatches 0
                                                                                   Mismatches
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                                                                                                                        Length 762;
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APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/185,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-38
PRIOR PR
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; TYPE: DNA
; ORGANISM: Human
; ORGANISM: Human
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US-10-027-632-3535
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US-10-027-632-3536
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PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-08-09
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Publication No. US20020198371A1
GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
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EQ ID NO 3535
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CURRENT FILING DATE: 2002-04-30
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; TYPE: DNA
; ORGANISM: Human
US-10-027-632-3536
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Query Match
32.1%; Score 17; DB 13; Length 762;
Best Local Similarity 100.0%; Pred. No. 8.9;
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps

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Qy 12 AAATCTGAATCAACCAG 28
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Db 474 AAATCTGAATCAACCAG 490

Search completed: December 5, 2004, 08:04:09 Job time : 235 secs

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Result
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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         seq length: 0
seq length: 2000000000
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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Copyright (c) 1993 - 2004 Compugen Ltd.
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    US-08-875-233-5
US-08-875-233-14
US-08-875-233-3
US-08-8-85-13-14
US-08-488-031-14
US-08-488-027-14
US-08-488-027-14
US-08-488-027-14
US-08-482-663-14
US-08-482-663-14
US-08-482-663-10-14
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Sequence 5, Appli
Sequence 14, Appli
Sequence 11, Appli
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INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 771 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                        NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5400
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                                                                                                                                                      MOLECULE TYPE: CI
HYPOTHETICAL: NO
ORIGINAL SOURCE:
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                                                                                              FEATURE:
                                                 NAME/KEY:
LOCATION:
    LOCATION: 3..660 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                              STRAIN: V-34
                                                                                                                                  ORGANISM:
                                                                     CDS
                                                                                                                                    Cucumber mosaic virus
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45.8 86.4 152 2 US-08-488-027-9 45.8 86.4 152 2 US-08-090-192-9 45.8 86.4 152 2 US-08-482-653-9 45.8 86.4 152 3 US-08-482-658-9 45.8 86.4 152 3 US-08-482-658-9 45.8 86.4 152 3 US-08-97-0349-9 45.8 86.4 152 5 PCT-US92-00277-9 45.8 86.4 152 5 PCT-US92-00277-9 45.8 86.4 154 1 US-08-090-193-13 45.8 86.4 154 2 US-08-488-031-13 45.8 86.4 154 2 US-08-488-027-13 45.8 86.4 154 2 US-08-482-658-13 45.8 86.4 154 3 US-08-492-658-13	<b>4</b> 5	44	43	42	41	40	39	38	37	36	35	34	ü	32	31	30	29	0
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	PCT-US92-00277-13	US-08-475-610-13	US-08-470-349-13	US-08-482-658-13		US-08-090-192-13	US-08-488-027-13	US-08-486-569-13	US-08-488-031-13	US-08-090-193-13	PCT-US92-00277-9	US-08-475-610-9	US-08-470-349-9	•	US-08-482-663-9	US-08-090-192-9	-08-488-027	-00-400-007-
	<ol> <li>Appl</li> </ol>	13, Appl	13, Appl	13, Appl	•	13, Appl	<ol> <li>Appl</li> </ol>	13, Appl	13, Appl	13, Appl	9, Appli	9, Appli	9, Appli	9, Appli	9, Appli	9, Appli	9, Appli	11 17 11

ALIGNMENTS

ERAL INFORMATION:  PPLICANT: Boeshore, Maury L  PPLICANT: McMaster, J. Russell  PPLICANT: Tricoli, David M  PPLICANT: Reynolds, John F  PPLICANT: Reynolds, John F  PPLICANT: Reynolds, John F  PPLICANT: Reynolds, John F  PPLICANT: Carney, Kim J  ITLE OF INVENTION: Cucumber Mosaic Virus  UMBER OF SEQUENCES: 15  ORRESPONDENCE ADDRESS:  ADDRESSEB: Rockey, Milnamow & Katz, Ltd.  STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700  CITY: Chicago  STATE: 11linois  COUNTRY: U.S.A.  ZIP: 60089  OMPUTER READABLE FORM:  MEDIUM TYPE: Floppy disk  COMPUTER: INM 'PC compatible  OPERATING SYSTEM: PC-DOS/MS-DOS  SOFTWARE: Patentin Release #1.0, Version #1.30  URRENT APPLICATION UNDERSE: US/OS/875, 233  PILING DATE: 26/JUN-1997
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/codon\_start= 3
/function= "ENCAPSIDATES VIRUS RNA"

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INFORMATION FOR SEQ ID NO: 1/
SEQUENCE CHARACTERISTICS:
LENGTH: 772 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                              Query Match
Best Local
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                                                                                                                                                                                                                  MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Cucumber Mosaic Virus
INDIVIDUAL ISOLATE: A35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:

APPLICATION UNBER: 08/68 875,233 FILING DATE: 26-UW-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: McMaster, J. Russell
APPLICANT: Tricoli, David M
APPLICANT: Reynolds, John F
APPLICANT: Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
TITLE OF INVENTION: Cucumber Mosaic Virus
NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
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NAME: Lisa V. Mueller
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OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
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STATE: Illinois
COUNTRY: U.S.A.
ZIP: 60089
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                     NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                  LOCATION:
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                                        4 CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                                              Similarity
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2 Prudential Plaza, 180 N. Stetson, Suite 4700
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                                                                                                                                                                                                                                                                                                                               linear
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                                                                         86.8%; Score 46; DB 3; Le 100.0%; Pred. No. 4.9e-09; tive 0; Mismatches 0;
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/gene= "CP"
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; Pred. No. 4.9
0; Mismatches
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US-08-090-193-14
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US-08-875-233-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 792 base pairs
TYPE: nucleic acid
CTRANSTRUCTS: 121
                                                                                                                                         Patent No.
                                                                                                                                                        Sequence 14,
                                                                                                                                                                                                                                                                                                                         Matches
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Best Local Similarity
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GENERAL INFORMATION:
APPLICANT: Boesho
APPLICANT: McMast
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                                                                                                                      GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
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MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICANT: Haseloff, James
APPLICANT: Haseloff, Andrea
APPLICANT: Brand, Andrea
APPLICANT: Perrimon, No. 5641673bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECUL: N
HYPOTHETICAL: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Plants Resistant to C Strains TITLE OF INVENTION: Cucumber Mosaic Virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY:
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FILING DATE: 26-JUN-1997
CLASSIFICATION: 800
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COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN: v-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Chicago
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5641673
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                                                                                                                                                                                                                                                                                                                       Conservative
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Tricoli, David M
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100.0%; Pred. No. 4.9e-(
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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APPLICANT: Haselo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 23-DEC-1993
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/FILING DATE: 16-JAN-1992
PRIOR APPLICATION NUMBER: US 07/64
PILING DATE: 17-JAN-1991
APPLICATION NUMBER: US 07/64
PILING DATE: 17-JAN-1991
ATTORNEY, AGENT INFORMATION:
NAME: Cimbala, Michele A.
REFERENCE/DOCKET NUMBER: 0609
REFIGERMACE/DOCKET NUMBER: 0609
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO:
                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,031
                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                            APPLICANT: Brand, Andrea
APPLICANT: Perrimon, No. 5849548bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                              14,
ING DATE: 07-JUN-1995
APPLICATION DATA:
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5849548
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                                                                                                                                                                                                                  Washington
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                                                                                                                                                                                                    D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              130 base pairs
                                                                                                                                                                                                                                     E: Sterne, Kessler, Goldstein & Fox 1100 New York Avenue, N.W., Suite 600
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                                                                                                                                                                               U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                        Haseloff, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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95.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 45.8; DB 1;
Pred. No. 3.6e-09;
0; Mismatches 2;
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                           FILING DATE: 07-UUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090
FILING DATE: 23-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US
FILING DATE: 16-UAN-1992
                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIAN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/486,569
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PILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
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LENGTH: 130 base pairs
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PRIOR APPLICATION DATA:
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080006/MAC/LBB
TELECOMMUNICATION INFORMATION:
                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 1100 New CITY: Washington
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STRANDEDNESS: both
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                                  NAME: Bugaisky, Lawrence B. REGISTRATION NUMBER: 35,086
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Perrimon, No. 5863774bert
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ER: 060
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TOPOLOGY: US-08-488-027-14
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                                  Matches
                                                  Query Match
Best Local Similarity
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INFORMATION FOR SEQ ID NO: :
                                                                                                                                                                                                            TELEFAX: (202) 371-25
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                        PILING DATE: 23-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/
FILING DATE: 16-JAN-1992
PRIOR APPLICATION NUMBER: US 07/64
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: BUGSLSKY, LEWENCE B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0669
TELECOMMUNICATION INFORMATION:
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                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 130 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PATENTIN Release #1.0, 1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,027
FILLING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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Local Similarity 95.9%;
es 47; Conservative
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GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
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VENTION: Cell Ablation Using Trans-Splicing Ribozymes
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                                                                                                                       linear
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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95.9%;
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Pred. No. 3.6e-09;
0: Mismatches 2;
                              Score 45.8; DB 2;
Pred. No. 3.6e-09;
0; Mismatches 2;
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                                                                  Length 130;
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                                                                                             GENERAL INFORMATION:
APPLICANT: Haselof
APPLICANT: Brand,
APPLICANT: Perrimo
                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                            Sequence 14,
                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                Patent No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 07/
FILING DATE: 17-VAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
REGISTRATION NUMBER: 33,85
REFERENCE/DOCKET NUMBER: 0
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APPLICATION NUMBER: PCT/
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Rel-Base #1.0, Version #1.25
CURRENT APPLICATION DATA:
                  APPLICANT: Perrimon, No. 588457777777 APPLICANT: Perrimon, No. 588457777777 Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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 STREET:
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                                                                                                                                                                1, Application US/08482663
5882907
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3: Sterne, Kessler, Goldstein & Fox
1100 New York Avenue, N.W., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                           130 base pairs
                                                                                         Brand, Andrea
Perrimon, No. 5882907bert
                                                                                                                                Haseloff, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       U.S.A.
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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Trans-Splicing Ribozymes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US 07/642,333
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Pred. No. 3
                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                             Gaps
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COMPUTER READABLE FORM: MEDIUM TYPE: Floppy

Floppy disk

20005

COMPUTER:

CITY: Wa STATE: C COUNTRY:

D.C.

U.S.A.

Washington

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US-08-482-658-14
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
             SOFTWARE: PATENTIN Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,658
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                         APPLICANT: Haseloff, James
APPLICANT: Brand, Andrea
APPLICANT: Perrimon, No. 6010904bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 23-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/U
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PAPLICATION NUMBER: US/08/482,0
FILING DATE: 07-UN-1995
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 08/090,193
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FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
                                                                                                                                                                                                                       STREET: 1100
CTTY: Washington
                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 0609.3080005/MAC/LBB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
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SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
                                                                                                                                                                                                                                                                                  ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  48 GAGTCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 96
                                                                                                                                                                                       20005
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                                                                                                                                                                                                                                                          E: Sterne, Kessler, Goldstein & Fox
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SYSTEM: PC-DOS/MS-DOS
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95.9%;
                                                                                        Release #1.0, Version #1.25
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08/090,193
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Pred. No. 3.6e-09;
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PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/US92/00277

PILING DATE: 16-JAN-1992

PRIOR APPLICATION NUMBER: US 07/642,330

PILING DATE: 17-JAN-1991

PILING DATE: 17-JAN-1991

ATTORNEY/AGENT INFORMATION:

NAME: Bugaisty, Lawrence B.

REGISTRATION NUMBER: 35,086

REFERENCE/DOCKET NUMBER: 0609.3080008/MAC/LBB

TELEPHONE: (202) 371-2540

INFORMATION FOR SEQ ID NO: 14:

SEQUENCE CHARACTERISTICS:

LENGTH: 130 base asize

LENGTH: 130 base asize
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                                                                                                                          APPLICATION NUMBER: 08/090,192
FILING DATE: 11-NOV-1993
APPLICATION NUMBER: PCT/ US 92/00278
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,333
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/470,349
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,192
                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Trans-Splicing Ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Haseloff, APPLICANT: Goodman,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
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                                                        TELEPHONE:
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X: (202) 371-2540
N FOR SEQ ID NO: CHARACTERISTICS:
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95.9%;
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Pred. No. 3.6e-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 3;
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US-08-475-610-14
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                                                                                                                                Query Match
Best Local Similarity
Matches 47; Conserv
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Best Local Similarity
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ZIP: 2005
ZIP: 2005
COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
PC-DOS/MS-DOS
TOTAL TOT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/090,193

FILING DATE: 23-DEC-1993

PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/00277

FILING DATE: 16-JAN-1992

PRIOR APPLICATION DATA:
APPLICATION DATA:
US 07/642,330

FILING DATE: 17-JAN-1991

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (202) 371-25
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                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 130 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERALIZATION DATA:
SOFTWARE: PatentIn Release #1..., .
CURRENT APPLICATION DATA:
CURRENT APPLICATION NUMBER: US/08/475,610
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APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid STRANDEDNESS: both TOPOLOGY: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080004/MAC/LBB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/01 FILING DATE: 07-JUN-1995
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                                                     GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
       GAGTCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 96
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                                                                                                                                            86.4%;
llarity 95.9%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Brand, Andrea
                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    86.4%;
95.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Release #1.0, Version #1.25
                                                                                                                                            Score 45.8; DB 3;
Pred. No. 3.6e-09;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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Pred. No. 3.6e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 3;
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                                                                                                                                                                                                                    Length 130;
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RESULT 14
PCT-US92-00278-14
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PCT-US92-00277-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; MOLECULE TYPE: DNA PCT-US92-00277-14
                                                                                                                                                                                 Sequence 14, Application PC/TUS9200278
GENERAL INFORMATION:
APPLICANT: Haseloff, James
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 51
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14, Application PC/TUS9200277 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (202)833-8716
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 130 bases
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 29,021
REFERENCE/DOCKET NUMBER: 0609.3496604
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jorge A
REGISTRATION NUMBER: 29,021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Brand, Andrea
APPLICANT: Perrimon, Norbert
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
                                                                                     STREET: La..
STREET: Washington
CITY: Washington
CTATE: DC
TTATE: DC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Haseloff, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            / Match 86.4%;
Local Similarity 95.9%;
nes 47; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: NUCLEIC ACID STRANDEDNESS: both TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: POFILING DATE: 19920116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
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                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE:
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                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                              48 GAGTCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                 20036
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1225 Connecticut Avenue, N.W., Suite 300
                                                                                                                                                  E: Sterne, Kessler, Goldstein & Fox 1225 Connecticut Avenue, N.W., Suite 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
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Pred. No. 3.6e-09;
0; Mismatches 2
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PC-DOS/MS-DOS n Release #1.0, Version #1.25

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RESULT 15
US-08-090-193-11
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SOFTWARE: Patentin Release #10, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/090,193
FILING DATE: 23-DEC-1993
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0609.3080001
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 11, Application US/08090193
Patent No. 5641673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 86.4%;
Best Local Similarity 95.9%;
Matches 47; Conservative
                                                                                                                                                                                                                                                                                                          COUNIA...
ZIP: 20005
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: IBM PC compatible
TYPEN: PC-DOS/MS-DOS
TYPEN: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 06/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)833-7533
TELEFAX: (202)833-8716
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Haselo:
APPLICANT: Brand,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: PCT/US92/00278
FILING DATE: 19920116
CLASSIFICATION: 435
CLASSIFICATION DATA:
APPLICATION NUMBER: US 07/642,333
APPLICATION NUMBER: US 07/642,333
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Brand, Andrea
APPLICANT: Perrimon, No. 5641673bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: PC-D
SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: NUCLEIC ACID
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Goldstein, Jorge A
REGISTRATION NUMBER: 29,021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
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1100 New York Avenue, Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     130 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Haseloff, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Linear
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Pred. No. 3.6e-09;
0; Mismatches 2;
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                                                                                                                             US-08-090-193-11
                                                             Query Match
Best Local Similarity
Matches 47; Conserv
                                                                                                                                                                                   TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 131 base pairs
                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: both
                                                                                                                                                                         TYPE:
49 GAGTCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 97
                  1 GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                              Conservative
                                                                          86.4%;
95.9%;
                                                             0
                                                           Score 45.8; DB 1;
Pred. No. 3.7e-09;
0; Mismatches 2;
                                                                                         Length 131;
                                                             Indels
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0

Search completed: December Job time : 53 secs 5, 2004, 06:38:56

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Title:
Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Database
Published Applications NA:*

1: /cgn2_6/ptcdata/2/pubpna/US07 PUBCOMB.seq:*

2: /cgn2_6/ptcdata/2/pubpna/US06_NEW PUB.seq:*

3: /cgn2_6/ptcdata/2/pubpna/US06_NEW PUB.seq:*

4: /cgn2_6/ptcdata/2/pubpna/US06_PUBCOMB.seq:*

5: /cgn2_6/ptcdata/2/pubpna/US08_NEW PUB.seq:*

6: /cgn2_6/ptcdata/2/pubpna/US08_NEW PUB.seq:*

7: /cgn2_6/ptcdata/2/pubpna/US08_NEW PUB.seq:*

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19: /cgn2_6/ptcdata/2/pubpna/US10E_PUBCOMB.seq:*

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20: /cgn2_6/ptcdata/2/pubpna/US10E_PUB.seq:*

20: /cgn2_6/ptcdata/2/pubpna/US10E_PUB.seq:*
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Gapop 10.0 , Gapext 1.0
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53
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 gatccatggacaaatctgaa.....gtcgtaaccgtcgacgagct 53
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

No.	Score	Match	Match Length DB	DB	ID	Description
1	46	86.8	771	13	US-10-011-033-5	Sequence 5, Appli
N	46	86.8	772	13	US-10-011-033-14	Sequence 14, Appl
u	46	86.8	792	13	US-10-011-033-3	Sequence 3, Appli
4	44.4	83.8	772	13	US-10-011-033-1	Sequence 1, Appli
ر ت	42.4	80.0	960	13	US-10-011-033-9	Sequence 9, Appli
6	24.8	46.8	983	13	US-10-011-033-11	GENERAL INFORMA
7	24	45.3	1068	8	US-08-961-527-373	Sequence 373, App
80	24	45.3	1068	16	US-10-158-844-373	Sequence 373, App
9	23.6	44.5	352	16	US-10-424-599-77125	Sequence 77125, A
10	23.2	43.8	1680	15	US-10-369-493-37626	Sequence 37626, A
11	22.8	43.0	642	10	US-09-905-666A-20	Sequence 20, Appl
12	22.8	43.0	766	17	US-10-437-963-47739	Sequence 47739, A

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444	44 4 3 3 8 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4		28 30 31	276	223	18 19 20	15
21.4		21.6 21.6 21.6		21.8	NNN	22.4	222.6
40.4	40.8 40.8 40.4	40.8 40.8 40.8 40.8			41.5		
746 1309 1386	4800 4800 401616 473 474	1554 1554 1922 2055 4800	255 255 735 1071 1488	58320 2256646	1180 1230 3708	1182 1266 3309400 1880	122859 536165 546
17 16	17 17 13	16 18 17	18 18 10 10		10 18		13
0-437-963 0-425-114 0-767-701	US-10-109-048-1141 US-10-628-525-4 US-10-087-192-1630 US-10-437-963-145 US-10-425-115-44987	US-10-425-114-35214 US-10-425-115-9544 US-10-425-115-953 US-10-425-115-953 US-10-437-963-11860 US-10-272-291-5	US-09-923-876-1370 US-09-923-876-1370 US-10-653-047-7425 US-10-425-114-35778 US-10-425-174-35778	US-10-767-795-1977 US-10-322-281-88 US-10-470-565-1	2 6	US-09-738-626-773 US-10-781-014-619 US-09-738-626-1 US-10-369-493-36410	US-10-087-192-37 US-10-087-192-37 US-09-939-964-1 US-10-027-632-200344 US-10-027-632-200344
	Sequence 1141, Ap GENERAL INFORMA Sequence 1630, Ap Sequence 145, App Sequence 44987, A	3521 9544 9553 1186		sequence 19//, Ap Sequence 88, Appl Sequence 1, Appli		7	Sequence 37, Appl Sequence 1, Appli Sequence 200344, Sequence 200344,

# ALIGNMENTS

STATE: CHICAGO STATE: ILLINOIS COUNTRY: U.S.A. ZIP: 60089 COMPUTER READABLE FORM: MEDIUM TYPE: Ploppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION NUMBER: US/10/011,033 FILING DATE: 13-No. US20020124286A1-2001 CLASSIFICATION LOTA: APPLICATION DATA: APPLICATION TUMBER: US/08/875,233 FILING DATE: 26-UN-1997 ATTORNEY/AGENT INFORMATION: TELECOMMUNICATION INFORMATION:	S/10011033 4286A1 4286A1 r, J. Russell r, David M s, John F Kim J F Kim J Cucumber Mosaic Virus S: 15 RESS: RESS: RESS: Ckey, Milnamow & Katz, Ltd. dential Plaza, 180 N. Stetson,
on #1.30	Strains of

TELEPHONE: 312-616-5400

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US-10-011-033-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 14, Application US/10011033 Publication No. US20020124286A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                  TELEFAX: 312-616-5460 INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                      APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION: <UNKNOWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 771 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ORIGINAL SOURCE:
ORGANISM: Cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                        ZIP: 60089

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Plants Resistant to C Strains Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /standard_name= "COAT PROTEIN" SEQUENCE DESCRIPTION: SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   h 86.8%; Score 46; DB 13;
Similarity 100.0%; Pred. No. 1.5e-08
46; Conservative 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Chicago
STATE: Illinois
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: 3..660
OTHER INFORMATION: /codon_start= 3
/function= "ENCAPSIDATES VIRUS RNA"
/product= "COAT PROTEIN"
/gene= "CP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN: V-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 312-616-5460
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              number=
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Carney, Kim J
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US-10-011-033-3
; Sequence 3, Application US/10011033
; Publication No. US20020124286A1
; GENERAL INFORMATION:
                                                                                                                                                                                     COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-NO. US20020124286A1-2001
CLASSIFICATION: <UNKNOWN-PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-UN-1997
ATTOREY/AGENT INFORMATION:
TELEPHONE: 11-616-5460
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-011-033-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 60089
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains of
Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORIGINAL SOURCE:
ORGANISM: Cucumber Mosaic Virus
INDIVIDUAL ISOLATE: A35
                    HYPOTHETICAL: NO ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: CDS
LOCATION: 3..660
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 15 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                   MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       46;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                                                                            LENGTH: 792 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: Illinois
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Rockey, Milnamow & Katz, Ltd
STREET: 2 Prudential Plaza, 180 N. Stet
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 46
                                                                                                          TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 772 base pairs
ORGANISM: CUCUMBER MOSAIC VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     86.8%; Score 46; DB 13; 100.0%; Pred. No. 1.5e-08; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 772;
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; NAME/KEY: CDS
; LOCATION: 3..660
; SEQUENCE DESCRIPTION: SEQ ID NO: 3:
US-10-011-033-3
                                                   ;
IAME/KEY: CDS
;
LOCATION: 3..660
;
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-10-011-033-1
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1, Application US/10011033 Publication No. US20020124286A1 GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 312-616-5460 INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIIM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-NO. US20020124286A1-2001
CLASSIFICATION APPLICATION DATA:
PRIOR APPLICATION UMBER: US/08/875,233
FILING DATE: 26-UN-1997
APPLICATION UMBER: US/08/875,233
FILING DATE: 26-UN-1997
APPLICATION THFORMATION:
TELEPHONE: 312-616-54400
TELEPHONE: 312-616-54400
                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 772 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ORIGINAL SOURCE:
ORGANISM: Cucumber Mosaic Virus
INDIVIDUAL ISOLATE: V-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains of
Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 15 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ch 86.8%; Score 46; DB 13; Similarity 100.0%; Pred. No. 1.5e-08; 46; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN: v-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            McMaster, J. Russell
Tricoli, David M
Reynolds, John F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Boeshore, Maury L
    83.8%;
97.8%;
    Score 44.4; DB 13;
Pred. No. 6.8e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 792;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                     Length 772;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>.</u>.
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RESULT 5
US-10-011-033-9
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밁
                                                                                                                                                        US-10-011-033-9
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                                                                           Matches
                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 312-616-5460 INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/11,033
FILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US/08/875,233
FILING DATE: 26-UN-1997
ATTORNEY, ACENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5400
                                                                                             Y Match 80.0%;
Local Similarity 97.7%;
                                                                                                                                                                                                                                                                       ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: CUCUMBER MOSAIC VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   McMaster, J. Russell
Tricoli, David M
Reynolds, John F
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains of
Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: cDNA HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                        SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Boeshore, Maury L
                                                                           43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          45;
                                      6 ATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 2 Pruden
CITY: Chicago
STATE: Illinois
ATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCATCGACG 44
                                                                                                                                                                                          NAME/KEY: CDS
LOCATION: 1...
                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 960 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGGCG 46
                                                                                                                                                                                                                                                   STRAIN: STRAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                           Conservative
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                                                                           0; Mismatches
                                                                                             Score 42.4; DB 13;
Pred. No. 4.8e-07;
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                                                                                                             Length 960;
                                                                           Indels
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                                                                           Gaps
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RESULT 6 US-10-011-033-11

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US-08-961-527-373
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Best Local Similarity
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                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                  APPLICANT: Charles F
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                          CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                         ADDRESSEE: Human Genome Sciences, Inc. STREET: 9410 Key West Avenue CITY: Rockville STATE: Maryland COUNTRY: USA
                                                                             COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOPTWARE: ASCII Text
FILING DATE:
CLASSIFICATION:
                                     APPLICATION NUMBER: US/08/961,527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/875,233
APPLING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: L15a V Mueller
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC Compatible
OCMPRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION: CUNKNOWN>
PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 60089
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Boeshore, Maury L
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                                                                                                                                                                                 20850
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATGGACANATCTGGATCTCCCAATGCTAGTAGAACCTCCCGGCG 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PAGES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 VOLUME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 312-616-5460
JOURNAL: J. Gen. Viro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                     Charles Kunsch

IVENTION: Streptococcus pneumoniae Polynucleotides and Sequences

EQUENCES: 391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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Tricoli, David M
                                                                                                                                     Diskette, 3.50 inch, 1.4Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1989
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Pred. No. 7.8;
O; Mismatches 12;
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RESULT 8
US-10-158-844-373
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                                                                             US-10-158-844-373
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 Matches
                     Best Local Similarity
                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (301) 309-8512 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                         NAME: Hyman, Mark J.
REGISTRATION NUMBER: 46,789
REFERENCE/DOCKET NUMBER: PB340P1D1
INFORMATION FOR SEQ ID NO: 373:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   / Match 45.3%;
Local Similarity 75.0%;
les 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Brookes, A. Anders
REGISTRATION NUMBER: 36,373
REFERENCE/DOCKET NUMBER: PB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: CD-R
COMPUTER: Dell Latitude Pentium
OPERATING SYSTEM: Windows 98
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/961,527
FILING DATE: 1997-10-30
APPLICATION NUMBER: US 60/029,960
FILING DATE: 1996-10-31
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/10/158,844
FILING DATE: 03-Jun-2002
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
CITY: Rockville
STATE: Maryland
                                                                                             TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 373:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Kunsch et al. TITLE OF INVENTION: Streptococcus pneumoniae Polynucleotides and Sequences NUMBER OF SEQUENCES: 391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                150 AATCTGCATCAACCAGTGCGTCACCTCAGCATCGACAAGC 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13 AATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACGAGC
 30;
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                                                                                                                                                                         ENGTH: 1068 base pairs
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D. US20040029118A1
Conservative
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                   45.3%; Score 24; DB 16; 75.0%; Pred. No. 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   373:
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 0; Mismatches
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                                       Length 1068;
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 Gaps
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RESULT 11
US-09-905-666A-20
; Sequence 20, App
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                                                                                                                                                                                                                                                                                   ; TYPE: DNA
; ORGANISM: Pseudomonas fluorescens
US-10-369-493-37626
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Clone ID: PAT_MRT3847_40658C.1
US-10-424-599-77125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-424-599-77125
                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 37626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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Best Local (
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SEQ ID NO 77125
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                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10(52052)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 352
TYPE: DNA
ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                     ENGTH: 1680
                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           155 ACTGTGAATGAACCATTGCTGGTGGTAAACGTATATGA 192
                                                                                                                                                                                                 34;
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                                                                                                                                        1 GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGAGC 52
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                                                                                                             GTTTCATGTGCGCATCTGAACGATGAATTGATGGCCGGCACCGAAGACGAGC 1072
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  Application US/09905666A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                   43.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 44.5%;
76.3%;
                                                                                                                                                                                            ; Score 23.2; D; Pred. No. 40; O; Mismatches
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Pred. No. 19;
0; Mismatches
                                                                                                                                                                                                                                          DB 15;
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                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                       Length 1680;
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FILE REFERENCE: 0184.310US
: CURRENT APPLICATION NUMBER: US/09/905,666A
: CURRENT FILING DATE: 2002-10-15
: PRIOR APPLICATION NUMBER: 60/217,954
PRIOR FILING DATE: 2000-07-13
: PRIOR APPLICATION NUMBER: 60/300,378
: PRIOR FILING DATE: 2001-06-21
: NUMBER OF SEQ ID NOS: 111
: SOFTWARE: Patentin Ver. 2.1
: SEQ ID NO 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       밁
문
                                                                                                                                                 ; OTHER INFORMATION: Clone ID: PAT_MRT4530_50479C.1 US-10-437-963-47739
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 47739, Application US/10437963
Publication No. US20040123343A1
                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                             SEQ ID NO 47739
LENGTH: 766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
                                                                                                                                                                                                                                                                                                                                       APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILS REFERENCE: 38-21(53221)B
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: NOVEL LIPASE GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: GIVER, LORRAINE J. APPLICANT: MINSHULL, JEREMY APPLICANT: VOGEL, KURT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Bacillus sp.
                                                                                                                                                                                                    ORGANISM: Oryza sativa
                                                                                                                                                                                                                           TYPE: DNA
                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 642
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300 AACCTTGTCCAACTCGGACTCTGCCAATCCTGGTCGTAGCCG 341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       436 CCAGGCACAGATCCAAATCAAAAATTCTTTACACATCCGTCTACAAGCT 485
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACGAGCT 53
                                   2 ATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cao,
                                                                                                                                                                                                                                                                                                                                                                                                                  Boukharov, Andrey A.
Barbazuk, Brad
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ilarity 66.0%;
Conservative
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                                                                                        43.0%;
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Pred. No. 46;
0; Mismatches
                                                                      Score 22.8; DI
Pred. No. 48;
0; Mismatches
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                                                                                                           DB 17;
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US-10-437-963-34792/c
Sequence 34792, Application US/10437963
Publication No. US20040123343A1
GENERAL INFORMATION:

APPLICANT: La Rosa, Thomas APPLICANT: Kovalic, David

RESULT 13

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RESULT 15
US-09-939-964-1
; Sequence 1, Application US/09939964
; Publication No. US20030054522A1
; GENERAL INFORMATION:
; APPLICANT: Rosenthal, Andre
; APPLICANT: Freiberg, Christoph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: MORFIS, David W.
APPLICANT: Engelhard, Eric K.
APPLICANT: Engelhard, Eric K.
TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: CANCER
FILE REPERENCE: 529452000122
CURRENT APPLICATION NUMBER: US/10/087,192
CURRENT FILING DATE: 2002-03-01
PRIOR APPLICATION NUMBER: US 09/747,377
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: US 09/798,586
PRIOR FILING DATE: 2001-03-02
NUMBER OF SEQ ID NOS: 2059
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US-10-087-192-37/c
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US-10-437-963-34792
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APPLICANT: Boukharov, Andrey A.
APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NO 34792
LENGTH: 2718
TYPE: DNA
TYPE: DNA
TYPE: DNA
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 37, Application US/10087192 Publication No. US20020182586A1
                                                                                                                                                                                                                                                                                                                                                                        Query Match
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NAME/KEY: misc_feature
LOCATION: (1)...(122859)
OTHER INFORMATION: n = A,T,C or G
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TYPE: DNA
ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                     Local Similarity 75.7
mes 28; Conservative
                                                                                                                                                                                                                          9102 AGCCTTGGACAAATCTTAATGAACGATTGGTGGAGGT 9066
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                                                                                                                                                                                                                                                                                                                  42.6%; Score 22.6; DB 13; Length 122859; 75.7%; Pred. No. 2.1e+02; tive 0; Mismatches 9; Indels 0;
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; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 536165
; TYPE: DNA
; ORGANISM: Rhizobium
US-09-939-964-1
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                                     Ś
                                                                                 Matches 31; Conservative
                                                                                                       Best Local
                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/939,964
CURRENT FILING DATE: 2001-08-27
PRIOR APPLICATION NUMBER: 09/214,808
PRIOR FILING DATE: 1999-06-22
NUMBER OF SEQ ID NOS: 1
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Perret, Xavier Philippe
APPLICANT: Broughton, William John
TITLE OF INVENTION: Genomic Sequence
TITLE OF INVENTION: Plasmid
                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE:
525007 GACGAACGTGGATCATGACGTGCTGATCGCAACGCTCGACCGGCT 525051
                                         9 GACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACGAGCT 53
                                                                                                     h 42.6%;
Similarity 68.9%;
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                                                                                 0; Mismatches
                                                                                                     Score 22.6; DB 10; Length 536165; Pred. No. 3e+02;
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                                                                                   0; Gaps
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Search completed: December 5, 2004, 07:06:42
Job time : 232 secs

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Maximum DB
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Match
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Ubiquitin
Coat prot
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Claim	Production ubiquitin m desired pro	WPI; 2000-4316 P-PSDB; AAY902	Fang R,	-alow)	11-DEC-1998	11-DEC-	22-JUN-2000	WO200036129-		CDS	Nicoti	Ubiquit: ds.	Ubiquitin	19-SEP-2000	AAA30858	RESULT 1 AAA30858 ID AAA3085			44		4 1	4 0	3 8 9 8	37		3 3 4	32 2	30	2 2 6 8	27	25	2 2 4 3	22
8; Page	tion of d tin monom d protein	000-43 ; AAYS	, Wu J	TRNI	1998,	-1998;	-2000.	36129-			ana ta	2.		-2000	58;	8 8 5			15	15	16	16	1 16	16	166	 	16	16	17	17	17	17 17	17
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42pp;	ired protein coding seque	37.	Chen X;	MOLECULAR A	98WO-SG000103	98WO-SG000103			/producta/partial	Location/Qualifie 647 /*tag= 7	٠		C-terminal	st entry)		DNA;			99	99	335199	13500	11475	11463	3118	639 762	542 542	35	334462	35026	1132	657 1128	657
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lish.	in plants or nce upstream			AGROBIOLOGY.	25 25				"Ubiquitin_monomer	lifiers		protein production; plant	fragment coding			BP.		ALI GNMENTS	ADJ65853 ADJ65854	ADG93175	ADC24703	AAZ45562	AAH78338	AAX34650	ABL09344	AAZ45575	AAS43396 AAI93810	AAT38246	ADC24763	AAA64890	AAC36474	AAQ80683 AAC45252	AAQ41742
	plant cells by linking a of the gene encoding the								_fragment"			cell; ubiquitin promoter;	sequence.		•		<i>F</i>	7	Adj65853 GLP-2 pep Adj65854 GLP-2 pep	Adg93176 Novel exp	H	Aaz45562 Nucleotid	₽ K	Aax34650 Starch br	Abblo9344 Drosophil	Aaz45575 Nucleotid	Aas43396 DNA encod Aai93810 Human pol	Aat38246 Y. pestis	Adc24763 Human wil	ADX95685 Human gen Aaa64890 Bordetell	Aac36474 Arabidops	Aag80683 CMV-SA co Aac45252 Arabidops	742 Cucumbe

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This sequence encodes the C-terminal fragment of a ubiquitin monomer. invention relates to a method for enhancing production of a desired

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Query Match
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Best Local (
                                                                                 Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based known CP sequences. The genes may be used for producing plants such as known country, peppers and tomatoes which are resistant to CMV grash, cucumber, peppers and tomatoes which are resistant to CMV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein in a plant or plant cell by inserting a nucleic acid (NA) encoding a ubiquitin monomer upstream of a NA encoding the desired protein, where the fusion construct encodes a fusion protein and expression is not controlled by the ubiquitin promoter. The invention also relates to a NA acid vector a NA vector able to transform a plant cell, that comprises NA encoding a fusion protein having a ubiquitin monomer linked to a protein of interest and further, where expression of the fusion construct is not under control of a ubiquitin promoter. The construct allows enhanced production of the desired protein in plants or
                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                               New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae, resistant to infection.
                                      Sequence 771 BP; 176 A; 199 C; 183 G; 213 T; 0 U; 0 Other;
                                                                                                                                                                               Claim 24; Fig 3; 80pp; English
                                                                                                                                                                                                                                                                                                                           Boeshore ML,
                                                                                                                                                                                                                                                                                                                                                                                        30-DEC-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cucumber mosaic virus; strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Coat protein; CP; cucumber mosaic virus; CMV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Coat protein
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02-DEC-1996
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                                                                       intection.
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DB; AAR98895.
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(first ent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of the V34 strain of cucumber mosaic virus
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                                                                       on 16-OCT-2003 to standardise OS field)
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           Score
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Pred. No. 6.1e-19;
; Mismatches 0;
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        Length 771;
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                                                                                                                                                                                                                                which are
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RESULT 4
AAQ10461
ID AAQ1
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AC AAQ1
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AAQ10461 standard;

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RESULT 3
AAT346655
ID AAT346657
AAT346657
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AAT34667
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AAT3467
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                                                                                                             Matches
                                                                                                                                                            Query Match
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Matches 46; Conserv
                                                                                                                                                                                                                                                                       Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tisses by PCR amplification of cDNA using primers based known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae,
                                                                                                                                                                                                                      Sequence 773 BP; 175 A; 200 C; 185 G; 213 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               plants, partic. of the resistant to infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Boeshore ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cucumber mosaic virus; strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT34665 standard; DNA; 773
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 13; Fig 2; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9621018-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Coat protein; CP; cucumber mosaic virus; CMV; V27; V33; V34; A35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT34665;
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                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1996-333993/33.
                                                                                                           46;
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                                                                                                                                     Similarity
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                            CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
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CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
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                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of the V33 strain of cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcmaster JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94US-00367789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                95WO-US007234
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            . 659
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                                                                                                           86.8%; Score 46; DB 2; L
100.0%; Pred. No. 3.4e-15;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  മ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tricoli DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ₽P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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hes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Reynolds JF,
                                                                                                                                                               Length 773
                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Carney KJ;
46
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                                                                                                           <u>,</u>
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                                                                                                           Gaps
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(revised)

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RESULT 5
AAQ10462
ID AAQ1
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AC AAQ1
XX
DT 27-A
DT 25-M
DT 16-A
XX
XX
XX
DE Caps
XX
XX
XX
CMV;
XX
CMV;
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Best Local S
Matches 45
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25-MAR-2003
16-APR-1991
                                                          27-AUG-2003
25-MAR-2003
16-APR-1991
                                                                                                                                                                                                                                                                      sequence introduced via an Agrobacterium tumefaciens intermediate are cultured as shoots in special media. The transformed melon plants are resistant to CMV. See also AAQ10462. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                               The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDS
                                                                                                                                                                                                                                                                                                                      to RNA 3 of the virulent strain FNY, isolated in New York on infected melons. The sequence includes a leader sequence, coding region and 3 coding region. The leader sequence is the same length as that of the Japanese Y strain of CMV. Transgenic melon plantlets containing the
                                                                                                                                                                                                                                                                                                                                                                                                     Transgenic melon plantlets prodn. from transformed shoots - by two-stage culture on specific medium for new transformed plants etc., esp. resistant to cucumber mosaic virus.
  Cucumber mosaic virus
                                      Capsid
                                                                                                                                                                                                                                                    Sequence 976 BP;
                                                                                                                                                                                                                                                                                                                                                                                  Claim 15; Page 17; 44pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-AUG-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-AUG-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EP412912-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CMV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Capsid protein
                   CMV; resistance; capsid protein;
                                                                                                 AAQ10462;
                                                                                                                     AAQ10462 standard; DNA; 1007
                                                                                                                                                                                                                                                                                                                                                                                                                                            P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1991-046027/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Deboth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-FEB-1991.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BIOC-) BIOCEM
                                                                                                                                                                                                                                                                                                                                                              gene was isolated from a plasmid (pUC18) containing DNA complementary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  resistance; capsid protein; Cucumis melo; ss
                                                                                                                                                                      74
                                       protein
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                                                                                                                                                                                                              45,
                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                 CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                                                                                                                      CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 118
                                                                                                                                                                                                           84.9%; Score 45; ilarity 100.0%; Pred. No. Conservative 0; Mismatc
                                                         (revised)
(revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 89FR-00010848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene of Cucumber Mosaic Virus strain
                                       gene of Cucumber Mosaic Virus strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90EP-00402282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
75. .731
                                                                                                                                                                                                                                                   224 A; 249 C; 230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag=  a
/product= "CMV strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Noe1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Z,
                                                                                                                     В₽
                                                                                                                                                                                                            Mismatches
                    Cucumis melo;
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                                                                                                                                                                                                                                                    273 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FNY capsid protein"
                                                                                                                                                                                                                               DB
                                                                                                                                                                                                                      1.2e-
                                                                                                                                                                                                                               <u>ب</u>
                                                                                                                                                                                                            0;
                                                                                                                                                                                                                              Length 976
                                                                                                                                                                                                                                                   U; 0 Other;
                                                                                                                                                                                                            Indels
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  FR2701960-A1.
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RRSULT 6
AAQ67
ID AAQ66
XX AAQ67
XX AAQ66
XX AAQ66
XX Caul
DT 25-M
DT 12--2
XX Caul
XX
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The gene was isolated from the virulent French strain 117F of CMV. Tomato plantlets at the 2-leaf stage were infected with CMV strain 117F. 15 days post-infection, the virus was purified from the infected leaves and cDNA was synthesised from RNAs 1,2,3 and 4. After purification and selection by standard techniques, cDNA complementary to RNA 4 was found to encode the capsid protein. It was recloned into "Blue scribe" plasmids and sequenced. Transgenic melon plantlets containing the sequence introduced via an Agrobacterium tumefaciens intermediate are cultured as shoots in special media. The transformed melon plants are resistant to CMV. See also AAQ10461. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Transgenic culture on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CDS
                                                                                                                                                                                                                                                                                                                                                                                                                  Cauliflower
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2003
12-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ67395;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1007 BP; 227 A; 259 C; 247 G; 274 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-AUG-1989;
                                                                                                                                                                                                                                       Cauliflower
                                                                                                                                                                                                                                                                                                                                           Cauliflower mosaic virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   resistant to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-AUG-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-FEB-1991
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                                                                                                                                                                                                                                                                                                              polyribozyme;
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45; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard; DNA; 1007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            melon plantlets prodn. from transformed shoots - specific medium for new transformed plants etc.,
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                                                                                                                                                                                                                                           mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                               mosaic virus capsid protein coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
(first en
                                                                                                                                                                                                                                                                                                              inactivate;
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/product=
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                                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ß
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         84.9%;
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                                                                                                                                                                                                                                                                                                           CMV; capsid protein; coat protein; inactivation; resistance; crop protection;
                                                                   "Capsid protein."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "CMV strain 117F capsid protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 2; Le
1.2e-14;
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                                                                                                                                                                                                                                                                                                                                                                                                                  sequence
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RESULT 7
AAT19545/
ID AAT19
XX AAT9
AC AAT9
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Best Local S
Matches 45
                                                                                                                                                                                                                                                        Transgenic plant; virus resistance; disease coat protein; antisense gene; ss.
                                                                                                                                                                                                                                                                                                                   25-MAR-2003
21-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The RNA encoding the capsid protein of cauliflower mosaic virus can be targetted by a nucleic acid sequence called a "polyribozyme". The polyribozyme has endoribonuclease activity and is able to inactivate the gene encoding the viral capsid protein. The polyribozyme comprises several catalytic regions derived from ribozymes and confers complete resistance to virus. See AAQ67391-94. (Updated on 25-MAR-2003 to correct
       07-MAY-1997;
                                                                                                    promoter
                                                                                                                                    sig_peptide
                                                                                                                                                                                                                                  Cucumber mosaic virus.
                                                                                                                                                                                                                                                                                           Cucumber mosaic virus RNA-3 cDNA fragment.
                                                                                                                                                                                                                                                                                                                                                      AAT99545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1007 BP; 227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New polyribozyme contg. several catalytic regions in complementary sequence - can inactivate gene for viral capsid protein, esp. for p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1994-281767/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-FEB-1993;
                              12-NOV-1997
                                                      EP806481-A2
                                                                                                                                                                                                3'UTR
                                                                                                                                                                                                                                                                                                                                                                              AAT99545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Fig 2; 67pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lenee P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-FEB-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-SEP-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GENE-) GENE SHEARS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            new virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                      CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Perez P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                 (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           resistant transgenic plants, also DNA sequence encoding
       97EP-00201379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93FR-00002269
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  93FR-00002269.
                                                                                                                                    /product= "coat
956. .1029
                                                                                                                                                                                  /*tag=
                                                                                                                                                           /*tag=
                                                                                                                                                                                                          Location/Qualifiers
                                                                                                  note= "coat protein leader sequence" 030. .1066
                                                                                                                          *tag=
                                                                                                                                                                         00. .958
                                                                                                                                                                                                                                                                                                                                                                            CDNA;
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; 259 C; 247 G;
                                                                           "F sequence
                                                                                                                                                                                                                                                                                                                                                                              1066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   b; Score 45; DB
b; Pred. No. 1.2
0; Mismatches
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                                                                                                                                                 protein"
                                                                            of sub-genomic promoter"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 274 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                    resistance;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                    RNA virus; CMV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           have the antisense gene construct integrated into their genomes. They are resistant to viral infection. In particular, they are resistant to CMV. Within the antisense gene construct, the interfering activity of the antisense foomain of the sub-genomic promoter, associated with the antisense activity performed by the coat protein gene, allows production of plants having 100% resistance to CMV. (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the subgenomic promoter of viral RNA, the coat protein gene, its leader sequence, and the tRNA-like 3'-terminal region of RNA-3. It was prepared by amplifying a fragment of CMV cDNA-3 from clone pCR-CMVIRNA3 by PCR (see also AAT99548-49). The gene construct is introduced into a vector containing a promoter active in plant cells in antisense orientation relative to the promoter. A claimed recombinant vector comprises the plant promoter, the antisense gene construct and a terminator which is functional in the plant. In addition to CMV, viral RNA may also be used from tobacco mosaic virus and potato virus. Claimed transgenic plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Preparing transgenic plants resistant anti:sense gene constructs containing from cucumber mosaic virus.
                                          Cellini F,
                                                                                                       09-MAY-1996;
                                                                                                                                     09-MAY-1997;
                                                                                                                                                                                                                              Cucumber mosaic virus
                                                                                                                                                                                                                                                         Transgenic plant; RNA virus; antisense construct; cucumber mosaic virus; CMV; promoter; coat protein gene; infection; RNA-3; ss.
                                                                                                                                                                                                                                                                                                       Cucumber mosaic virus (CMV) RNA-3 gene cDNA clone fragment
                                                                                                                                                                                                                                                                                                                                     26-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                     AAZ07505
                                                                                                                                                                                                                                                                                                                                                                                                 AAZ07505 standard; cDNA; 1067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1066 BP; 299 A; 251 C;
                                                                                                                                                                   28-SEP-1999
                                                                                                                                                                                                US5959181-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This cDNA clone of cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Page 10; 18pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1997-538620/50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-MAY-1996;
                                                                         (META-) METAPONTUM AGROBIOS SCRL
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7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                            Grieco PD
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                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                       96IT-MI000927.
                                                                                                                                     97US-00854170
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                   BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                273 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   to RNA virus infection
the viral coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                243 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 2; Lo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CMV) RNA-3 includes
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WPI; 1997-538620/50

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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to preparing transgenic plants resistant to RNA virus induced infections that comprises integrating an antisense gene construct into the plant genome. The construct comprises: (a) an F domain of a subgenomic promoter of cucumber mosaic virus (CWV); (b) downstream from the subgenomic promoter, a leader sequence of a coat protein gene of CWV; (c) downstream from the leader sequence, a gene encoding a CMV coat protein; and (d) downstream from the gene, a 3'-terminal region of a CMV coat protein gene. The method is useful for producing plants which are resistant to infection by RNA based viruses. The gene construct gives higher levels of resistance compared to antisense constructs which are capable of complementing with different domains of genomic RNA of CMV. The present sequence represents the fragment of cDNA clone of RNA-3 of CMV. This forms the antisense construct of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Preparing transgenic plants resistant to RNA virus infection - using anti:sense gene constructs containing the viral coat protein gene, e.g. from cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                EP279433-A
                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                       Cucumber mosaic virus; plant viral resistance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence contg.
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12-NOV-1990
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                                                                                    P-PSDB;
                                                                                                                           Furusawa I,
                                                                                                                                                                                                20-FEB-1987;
25-FEB-1987;
                                                                                                                                                                                                                                          18-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAN81111;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAN81111 standard; DNA; 1379
                                                                                                                                                       (TAKE )
                                                                                                                                                                                  18-FEB-1988;
                                        coding for the coat protein of cucumber mosaic virus strain Y producing plants resistant to cucumber mosaic virus infection.
                                                                                                 1988-236708/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2; Fig 1; 15pp; English.
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                                                                                    AAP80509.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                           Onda H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first en
                                                                                                                                                       CHEM IND LTD.
                                                                                                                                                                                  87JP-00043443.
88JP-00035809.
                                                                                                                                                                                                              87JP-00038288
                                                                                                                                                                                                                                          88EP-00102322
                                                                                                                                                                                                                                                                                                                                                                                                                                                     CMV strain Y coat protein gene
                                                                                                                                                                                                                                                                                                                          /label= cucumber mosaic virus-Y coat protein
                                                                                                                                                                                                                                                                                                                                                                  location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       297 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          84.9%; ST
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                       .1074
                                                                                                                           Komiya
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 45;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       244 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 2; L
1.2e-14;
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RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This DNA is produced on screening of a plasmid library and is used to transform plant cells which subsequently produce the coat prot- ein of cucumber mosaic virus (CMV) strain Y. This protein is not synthesised in natural plant cells. The resistance to CMV infect- ion, provided by the DNA, is shown in plant cells and redifferent- iated plant bodies. See also AAN81110. (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                        P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                         virus resistance; disease resistance; transgenic plant; cross hypersensitive response; crop improvement; tomato; Lycopersicon esculentum; tobacco; Nicotiana tabacum; RNase Tl;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2003
20-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1379 BP; 306 A; 342 C; 333 G;
                                                                                                           eliciting agent.
                                                                                                                      DNA constructs for imparting pathogen, espendes RNA that interacts with viral RNA
                                                                                                                                                                                          De Haan PT;
                                                                                                                                                                                                               (SANO)
                                                                                                                                                                                                                                                                                03-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                           Cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cucumber mosaic virus; tomato mosaic virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cucumber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ76107 standard; DNA; 1696 BP
                                                                                                                                                                                                                                                           04-JUN-1993;
                                                                                                                                                                                                                                                                                                                              WO9429464-A1
                                                                                                                                                                                                                                                                                                                                                                                                                               cell inhibitory protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                  1995-036490/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      417
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                                                                                                                                                             SANDOZ LTD.
SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN
                                                                                                                                                         AAR67754.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mosaic virus RNA-3 chimera encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
                                                                                                                                                                                                                                                          93GB-00011593
                                                                                                                                                                                                                                                                                 94WO-EP001817
                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                  /*tag= a
/product= "RNase Tl'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      84.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                TOMV; CMV; 88.
                                                                                                                                                                                                                 VERW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 °,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 45;
Pred. No.
                                                                                                                                                                                                                GES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              398 T; 0 U; 0 Other;
                                                                                                                       esp. virus, ro
NA polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.2e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           minus-strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 1379;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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                                                                                                                      resistance to
se to generate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                        plants
an
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A chimeric cucumber mosaic virus RNA-3 (given in AAQ76107) has the coat protein gene replaced by a gene encoding a cell inhibitory protein, RNase T1, having the sequence given in AAR67754. The construct elicits a minussense RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus-resistance on a host plant, e.g. tobacco, tomato. (Updated on 25-MAR-2003 to correct PN field.)

Claim 4; Page 31-32; 50pp;

English.

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RESULT 11

AAQ76106
ID AAQ76
AC AAQ76
AC AAQ76
AC AAQ76
AC Cucun
KW Cucun
KW Lycop
KW ToMV,
XX Cucur
XX TOMV,
XX Cucur
X
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                            Query Match
Best Local
     Matches
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                                                                                                                                                                         A chimeric cucumber mosaic virus RNA-3 (given in AAQ76106) codes for the coat protein (CP) of tomato mosaic virus (AAR67752) as well as its own CP (AAR67753). The construct elicits minus-sense RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus-resistance on a host plant, e.g. tobacco, tomato. (Updated on 25-MAR-2003)
                                                                                                                                                                                                                                                                                                                                                                                                   DNA constructs for imparting pathogen, esp. virus, resistance to encodes RNA that interacts with viral RNA polymerase to generate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1995-036490/05.
P-PSDB; AAR67752, AAR67753.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       virus resistance; disease resistance; transgenic plant; cross protecti hypersensitive response; crop improvement; tomato; Lycopersicon esculentum; tobacco; Nicotiana tabacum; coat protein; CP;
                                                                                                    Sequence 1860 BP; 448 A; 429 C; 431 G;
                                                                                                                                                    resistance on a host plant, e.g. to correct PN field.)
                                                                                                                                                                                                                                                                                                                           Claim 3; Page 27-28; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                          eliciting agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        De Haan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SANO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-JUN-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9429464-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cucumber mosaic virus RNA-3 chimera encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-2003
20-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ76106 standard; DNA; 1860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cucumber mosaic virus; tomato mosaic virus; minus-strand
     45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        σı
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SANDOZ LTD.
SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN VERW
                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1696 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 776
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       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
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895. .1550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                *tag=
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100.0%; Pred. No.
84.9%; Score 45;
100.0%; Pred. No.
cive 0; Mismatcl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 413 C; 389 G; 507 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "CMV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         coat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GES
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       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 coat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein"
                                                                                                    552 T; 0 U; 0 Other;
                                                   DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 2;
                              1.2e-14;
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                                                   Length 1860;
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     Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA;
     0
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  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                      plants
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RESULT 12
AAQ76108
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                                                         Query Match
Best Local Similarity
                                              Matches
                                                                                                                 A chimeric cucumber mosaic virus RNA-3 (given in AAQ76108) has the coat protein gene replaced by a gene encoding an elicitor, TOMV P30, having the sequence given in AAR67755. The construct elicits a minus-sense RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus-resistance on a host plant, e.g. tobacco, tomato. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                        DNA constructs for imparting pathogen, esp. virus, resistance to encodes RNA that interacts with viral RNA polymerase to generate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cucumber mosaic virus; tomato mosaic virus; minus-strand RNA; virus resistance; disease resistance; transgenic plant; cross protection; hypersensitive response; crop improvement; tomato; Lycopersicon esculentum; tobacco; Nicotiana tabacum; P30; elicitor; TOMV;
                                                                                            Sequence 2173 BP; 561 A; 438 C; 532 G; 642 T; 0 U; 0 Other;
                                                                                                                                                                                                                             eliciting
                                                                                                                                                                                                                                                                                                                                      (SANO)
(SANO)
                                                                                                                                                                                                                                                                                                                                                                                                              03-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cucumber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ76108
                                                                                                                                                                                                        Claim
                                                                                                                                                                                                                                                                                                                 De Haan
                                                                                                                                                                                                                                                                                                                                                                                       04-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                     22-DEC-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9429464-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cucumber mosaic
1209 CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 1253
                                                                                                                                                                                                                                                                                        1995-036490/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          968
                                                                                                                                                                                                     5; Page 34-35; 50pp; English.
                      IJ
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                                                                                                                                                                                                                                                                                                                                      SANDOZ LTD.
SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN VERW
                                                                                                                                                                                                                                                                                                                 PT;
                                                                                                                                                                                                                                                                              AAR67755
            CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard; DNA;
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                                                                                                                                                                                                                             agent.
                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                              94WO-EP001817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BULIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           virus RNA-3 chimera encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /product= "ToMV P30 elicitor"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              *tag=
                                                        84.9%; Score 45;
100.0%; Pred. No.
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                                              <u>,</u>
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                                              Mismatches
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                                                         DB 2; L
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                                              0
                                                                   Length 2173
                                              Indels
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                                              <u>.</u>.
                                              Gaps
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RESULT 13 AAT34664

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AAT17259
ID AAT1
XX
AC AAT1
XX
AC AAT1
DT 16-C
DT 02-D
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COat
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KW Coat
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COS
CCCC
XX
CCC
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                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based (known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-OCT-2003
02-DEC-1996
               Cucumber mosaic virus; strain
                                             Coat protein; CP; resistance; ss.
                                                                                              Coat protein
                                                                                                                             16-OCT-2003
02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; Fig 1; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      resistance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Coat protein; CP; cucumber mosaic virus; CMV; V27; V33; V34; A35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Coat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT34664;
                                                                                                                                                                               AAT17259
                                                                                                                                                                                                              AAT17259 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1996-333993/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Boeshore ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-JUL-1996.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT34664 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           resistant to infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ASGR-)
                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein of the V27 strain of
                                                                                                                                                                                                                                                                                                                                                            43;
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                                                                                                                                                                                                                                                                                                                            CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCG 46
                                                                                                                                                                                                                                                                                                                                                                                                                         772 BP; 177 A; 202 C; 182 G; 211 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                             CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88
                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                              of the A35 strain of cucumber mosaic virus
                                                                                                                              (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcmaster JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94US-00367789
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/*tag=
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                                                              cucumber
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                                                                                                                              entry)
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100.0%;
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                                                                mosaic virus;
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                                                                                                                                                                                                                                                                                                                                                                           Score 43;
Pred. No.
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                A35
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                                                                                                                                                                                                                                                                                                                                                                           1.5e-13;
                                                                CMV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Reynolds JF,
                                                                V27;
                                                                                                                                                                                                                                                                                                                                                                                         Length 772;
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                                                                V34;
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ABL58209
ID ABL58
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Best Local
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                                                                                                                                                                                                              CDS
               (RURA-) RURAL
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Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based c known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CDS
                                                                                                                                                                                                                                                                                                                                                                                                 ABL58209;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Fig 8; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated cucumber mosaic virus coat protein DNA - used to produce plants, partic. of the family Cucurbitaceae or Solanaceae, which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-333993/33.
P-PSDB; AAR93803.
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17-MAR-1998;
                                                                                                                                                                                                                                   Agrobacterium
                                                                                                                                                                                                                                                                                     Agrobacterium
                                                                                                                                                                                                                                                                                                                          Agrobacterium
                                                                                                                                                                                                                                                                                                                                                              25-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                   ABL58209
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                                                                      05-OCT-1999
                                                                                                          KR99075101-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      resistant to infection.
                                   17-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
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                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                     tumefaciens
                                                                                                                                                                                                                                                                                       tumefaciens; anti-cucumber mosaic virus;
                                                                                                                                                                                                                                                                                                                       tumefaciens coding sequence
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 98KR-00009096
                                   98KR-00009096
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                                                                                                                                           product=
                                                                                                                                                          *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA; 894
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100.0%; Pred. No.
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                                                                                                                                           "Agrobacterium tumefaciens
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. 1.5e-13;
ches 0;
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                                                                                                                                           protein"
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                                                                                                                                                                                                                                                                                         transgenic tomato;
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DEV ADMINISTRATION

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PI Sohn SH, Lee SW, Park JS, Hwang YS;

XX

DR WPT; 2000-577926/54.

DR P-PSDB; ABB80503.

XX

PT Anti-cucumber mosaic virus tomato.

XX

CC Disclosure; Page 4; 6pp; Korean.

XX

CC This sequence represents a coding sequence from Agrobacterium tumefaciens cused within the scope of the invention. The invention relates to an anti-

CC cucumber mosaic virus tomato

XX

Sequence 894 BP; 188 A; 228 C; 216 G; 262 T; 0 U; 0 Other;

Query Match

Best Local Similarity 100.0%; Pred. No. 7.6e-11;

Matches 38; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy

12 AAATCTGAATCAACCAGTGCTGGTTGGTAACCGTCGACG 49

11 AAATCTGAATCAACCAGTGCTGGTTGGTAACCGTCGACG 148

Search completed: December 5, 2004, 07:10:27

Job time: 223 secs
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70.735 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

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ADE58906 ADD45534 ADE61258 ADE58902	AAM12356 AAU23534 ABU11773 AAM39480 ABB61652	AAG05253 AAG00746 ABO57865 ABC57865	AAR 98894 AAR 98895 AAR 98895 ABB 80503 ADD 61973 AAD 7717	AAY90255 AAR10652 AAR10653 AAR57968 AAR67753 AAR688803	ID
Ade58906 Human Pro Add45534 Human Pro Ade61258 Human Pro Ade58902 Human Pro	Aamqız66 Human poı Aau23534 Novel hum Abu11773 Human MDD Aam39480 Human pol Abb61652 Drosophil	Arabic Human Human Human	Adrysos Coat prot Aar98894 Coat prot Aar98895 Coat prot Abb80503 Agrobacte Ado61973 Transcrip Aar37212 Croumber	Ubique Capsi Capsi Capsi Cauli	

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53.6	53.6	53.6	53.6	53.6	55.1	55.1	55.1	55.1	55.1	55.1	55.1	55.1	55.1	55.1	55.1	55.1	55.1	55.1	55.1
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ABM61147	AAU64628	AAU66349	AAG26614	ABP03145	ABP76680	AAG38480	AAG38481	AAG38482	AAY32080	ADP43242	ADP43243	ADP43232	ADC08287	ADC07876	AAY67287	AAR22239	ADO57439	ABM38071	AAU41552
Abm61147	Aau64628	Aau66349	Aag26614	Abp03145	Abp76680	Aag38480	Aag38481	Aag38482	Aay32080	Adp43242	Adp43243	Adp43232	Adc08287	Adc07876	Aay67287	Aar22239	Ado57439	Abm38071	Aau41552
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## ALIGNMENTS

promoter.

Production of desired protein in plants or plant cells ubiquitin monomer coding sequence upstream of the gene desired protein. by linking a encoding the

This sequence represents the C-terminal fragment of a ubiquitin monomer. The invention relates to a method for enhancing production of a desired protein in a plant or plant cell by inserting a nucleic acid (NA) encoding a ubiquitin monomer upstream of a NA encoding the desired protein, where the fusion construct encodes a fusion protein and expression is not controlled by the ubiquitin promoter. The invention expression is not controlled by the ubiquitin promoter. The invention cell, that comprises NA encoding a fusion protein having a ubiquitin monomer linked to a protein of interest and further, where expression of the fusion construct is not under control of a ubiquitin promoter. The construct allows enhanced production of the desired protein in plants or

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RESULT 3
AAR10653
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Matches 14
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Best Local (
                                                                                                                                                                                                                                                                                       The gene encoding this protein was isolated from a plasmid (pUC18) containing DNA complementary to RNA 3 of the virulent strain FNY, isolated in New York on infected melons. Transgenic melon plantlets containing the nucleotide sequence introduced via an Agrobacterium tumefaciens intermediate are cultured as shoots in special media. The transformed melon plants are resistant to CMV. See also AAQ10462. (Updated on 25-MAR-2003 to correct PI field.) (Updated on 27-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Transgenic melon plantlets prodn. from transformed shoots - by two-stage culture on specific medium for new transformed plants etc., esp. resistant to cucumber mosaic virus.
             AAR10653;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                              AAR10653 standard;
                                                                                                                                                                                                                                                             Sequence 218 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB; AAQ10461.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Deboth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cucumber mosaic virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Capsid protein of Cucumber Mosaic Virus strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-AUG-2003
25-MAR-2003
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                                                                                                                                                                                            1 Similarity
14; Conser
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                                                                                                                             MDKSESTSAGRNRR 14
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                                                                                                                                                                                              Conservative
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                                            protein; 218 AA.
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Pred. No. 3.8e-05;
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Pred. No.
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                                                                                                                                                                                            . 0.00071;
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AAR57968
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25-MAR-2003
16-APR-1991
                                                                                                                                                                                                                                            25-MAR-2003
20-APR-1995
                             26-FEB-1993;
                                                                                                                           Cauliflower mosaic virus.
                                                                                                                                                                                                           Cauliflower mosaic virus capsid protein
                                                                                                                                                                                                                                                                                            AAR57968;
                                                                                                                                                                                                                                                                                                                            AAR57968 standard; protein; 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 218 AA;
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                                                             02-SEP-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-FEB-1991
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(first entry)
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 69;
Pred. No.
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The gene encoding this protein was isolated from the virulent French strain 117F of CMV. Transgenic melon plantlets containing the coding sequence (introduced via an Agrobacterium tumefaciens intermediate) ar cultured as shoots in special media. The transformed melon plants are resistant to CMV. See also AAQ10461. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PA field.)
Cauliflower mosaic virus; CMV; capsid protein; coat protein; polyribozyme; inactivate; inactivation; resistance; crop pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Transgenic melon plantlets prodn. from transforculture on specific medium for new transformed resistant to cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CMV; strain 117F; resistance; capsid protein; Cucumis melo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             from transformed shoots -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.00071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2;
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protection.
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RESULT 5
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Best Local S
Matches 14
                                                                                               (SANO)
(SANO)
(SANO)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polyribozyme contg. several catalytic regions in complementary sequence - can inactivate gene for viral capsid protein, esp. for of new virus resistant transgenic plants, also DNA sequence encodi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                                                                                                                                                                              virus resistance; disease resistance; transgenic plant; cross protection; hypersensitive response; crop improvement; tomato; Lycopersicon esculentum; tobacco; Nicotiana tabacum; coat protein; CP;
         DNA constructs for imparting pathogen, encodes RNA that interacts with viral
                                                                                                                                                                                                                                                                                                                       CMV coat protein
                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
20-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                              AAR67753;
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                                                                          De Haan
                                                                                                                                            04-JUN-1993;
                                                                                                                                                                03-JUN-1994;
                                                                                                                                                                                       22-DEC-1994.
                                                                                                                                                                                                            WO9429464-A1
                                                                                                                                                                                                                                 Cucumber mosaic
                                                                                                                                                                                                                                                                                                 Cucumber mosaic virus; tomato mosaic virus; minus-strand RNA;
                                                                                                                                                                                                                                                                                                                                                                                                    AAR67753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 218
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                                                                                                                                                                                                                                                       TOMV;
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                                          1995-036490/05.
)B; AAQ76106.
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                                                                                               SANDOZ LTD.
SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          MDKSESTSAGRNRR 14
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(first en
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         esp. virus, resistance to RNA polymerase to generate
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encodes RNA that eliciting agent.

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MDKSESTSAGRNRR

Matches Query Match Best Local :

14;

Conservative

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Mismatches

Similarity

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Score 69; Pred. No.

DB 2; LC 0.00071; 0;

Length 218;

Indels

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                                                                                                                                                                                                                                                                                                                                                                                    WPI;
  Sequence
                                          Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based cknown CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                               New isolated cucumber mosaic virus coat protein plants, partic. of the family Cucurbitaceae or resistant to infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-OCT-2003
02-DEC-1996
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                                                                                                                                                                                                                Disclosure; Fig 8; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cucumber mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Coat protein; resistance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Coat protein
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DB; AAT17259.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 30; 50pp;
  AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of the A35 strain of cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Mcmaster
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    218
                                                                                                                                                                                                                                                                                                                                                                                                                                      Tricoli DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>,</u>
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 virus;
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. 0.00071;
. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Reynolds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CMV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 V27;
                                                                                                                                                                                                                                                                                        n DNA - used
Solanaceae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 218;
                                                                                                                                                                                                                                                                                                                                                                                                                                      JF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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RESULT 7
AAR98893
ID AAR9
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XX Coat
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ARPSHIT 8
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KW resi
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tisses by PCR amplification of cDNA using primers based on known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                  16-OCT-2003
02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Fig 1; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated cucumber mosaic virus coat protein plants, partic. of the family Cucurbitaceae or resistant to infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR98893;
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                Cucumber mosaic virus; strain V33.
                                                                                                                                                Coat protein
                                                                                                                                                                                                                                                                                    AAR98894;
                                                                                                                                                                                                                                                                                                                                  AAR98894 standard; protein; 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 218 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Boeshore ML,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cucumber mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 resistance.
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                                                                                            Coat protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT34664.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                of the V33 strain of cucumber mosaic virus.
                                                                                                                                                                                                  (revised)
(first entry)
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcmaster JR,
                                                                                            CP; cucumber mosaic virus;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 69;
Pred. No.
                                                                                                                                                                                                                                                                                                                                       8
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                                                                                               CMV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CMV; V27; V33; V34; A35;
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                                                                                          V27; V33; V34; A35;
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Solanaceae,
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ID AARPS
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DT 16-0
DT 02-D
XX COAT
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Best Local
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N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based (known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae,
                                                                         WPI; 1996-333993/33.
N-PSDB; AAT34666.
                                                                                                                                                                                                                                                                                             07-JUN-1995;
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02-DEC-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated cucumber mosaic virus plants, partic. of the family Cucur resistant to infection.
                                                                                                                                                                                                                                                30-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                11-JUL-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Cucumber mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Coat protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Coat protein of the V34 strain of cucumber mosaic virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR98895 standard; protein; 218
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-DEC-1994;
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                                                                                                                                                                                                  (ASGR-) ASGROW
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DB; AAT34665.
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ilarity 100.0%;
Conservative (
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(first entry)
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                                                                                                                                                  Mcmaster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mcmaster JR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                             virus; strain V34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cucumber mosaic virus; CMV; V27; V33; V34;
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                                                                                                                                                  Tricoli DM,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virus coat protein DNA - used Cucurbitaceae or Solanaceae,
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                                                                                                                                                  Reynolds JF,
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                                                                                                                                                  Carney
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  to produce which are
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RESULT 11
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                                                                                                                                                                                                                           Query Match
Best Local S
Matches 13
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Best Local S
Matches 14
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                                                                                                                                                                                                                                                                                                                           Sequence 218
                                                                                                                                                                                                                                                                                                                                                                                                    This sequence represents a protein sequence from Agrobacterium tumefaciens used within the scope of the invention. The invention relates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-OCT-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 218
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anti-cucumber mosaic virus tomato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-577926/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Agrobacterium
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 resistant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 MDKSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14;
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                                                                                                                          MAKSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lee
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                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98KR-00009096
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                                                                                                                                                                                                                                                                                                                                                                             mosaic virus tomato
                                                                                                                                                                                                                                                  92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 69; DB 2;
100.0%; Pred. No. 0.00071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               anti-cucumber mosaic virus; transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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                                                                                                                                                                                                                                                  Score 61; Pred. No. 0
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                                                                                                                                                                                                                           Mismatches
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0.018;
                                                                                                                                                                                                                                                                          Length 218
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ADO61973 standard; protein; 438

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CC germination in heat, increased tolerance to freezing conditions, concern controlled in heat, increased tolerance to disease, including concern conditions, increased tolerance to disease, including contentions, and contentions, increased tolerance to disease, including contentions, and contentions, increased tolerance to disease, including contentions, and contentions, and contentions, contention, conte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to novel plant transcription factor proteins (I) and nucleotide sequences (II) (ADO61534-ADO63778). The sequences can be used to produce transgenic plants, which overexpress (II), where the transgenic plant has an altered trait as compared to a non-transgenic plant or wild-type plant. The transgenic plant comprises an altered trait selected from increased tolerance to abiotic stress, increased tolerance to cold, increased germination in cold, increased tolerance to heat, increased increased germination in cold, increased tolerance to heat, increased
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New recombinant polynucleotide encoding transcription factor polypeptides, useful for producing transgenic plants with advantageous properties compared to a reference plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jiang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-SEP-2002; 2002US-0411837P
17-DEC-2002; 2002US-0434166P
24-APR-2003; 2003US-0465809P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plant; transcription factor; transgenic plant; abiotic stress tolerance; osmotic stress tolerance; cold tolerance; heat tolerance; low phosphate tolerance; fungal disease; low phosphate tolerance; fungal disease; glyphosate resistance; flowering; fertility; seed development.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 440; 510pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-SEP-2003; 2003WO-US030292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arabidopsis thaliana.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
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V, Dubell A
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7, Keddie
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Sherman |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      맖;
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AAR3721
AAR372
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AAR37212
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                                                                                  Matches
                                                                                                              Query Match
Best Local (
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                                                                                                                                                                                                                                          The 3' terminal 140-320 bases of the segmental genome RNA sequences of Bromovirus, Alfalfa mosaic virus and Cucumovirus (including CuMV) show a high degree of homology and this information was used to design a primer (see AAQ41743) complementary to part of the 3' terminal non coding region of the RNA3 (encodes the coat protein) of CuMV-Q which was used to prime synthsis of a cDNA sequence using RNA3 of CuMV-W as a template. The ds cDNA corresp. to RNA3 of CuMV-W was introduced into plasmid pUC19 and used to transform E. coli. Transformants contg. an insert of 800 bp were selected and sequenced. The CuMV-W coat protein gene sequence may be used to transform plants to produce transgenic plants with resistance to CuMV infection. Protected plants are puref. maize or from Curcurbitaceae or Solanaceae species, e.g. cucumber, melon, squash, tomato, potato or tobacco. (Updated on 24-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Protecting plants against cucumber mosaic virus infection - relates to e.g. isolated coat protein of virus strain and isolated DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-1992.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CuMV-W; transformants; transgenic plant; maize; Curcurbitaceae; Solanaceae; melon; squash; tomato; potato; tobacco.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-OCT-2003
26-AUG-1993
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                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-JUN-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cucumber mosaic virus-W coat protein.
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(UYCA-) UNIV OF CAPE TOWN.
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                                                                                  Similarity 9; Conservat
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MDKSESTSAGRNRR 14
                                                                                                                                                                                             218
                                                                                    Conservative
                                                                                                                                                                                             AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
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                                                                                                              60.9%;
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                                                                                                              Score 42; DB
Pred. No. 39;
                                                                                    Mismatches
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                                                                                                                                       DB 2;
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                                                                                                                                     Length 218;
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RESULT 13

AAG055553

AAG055553

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14-JUN-1999;
16-JUN-1999;
16-JUN-1999;
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20-MAY-1999;
21-MAY-1999;
24-MAY-1999;
25-MAY-1999;
27-MAY-1999;
27-MAY-1999;
01-JUN-1999;
03-JUN-1999;
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07-MAY-1999;
11-MAY-1999;
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05-MAR 1999
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21-APR-1999;
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  17-JUN-1999;
18-JUN-1999;
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07-JUN-1999;
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18-MAY-1999;
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99US-0129845P.
99US-0130077P.
99US-0130449P.
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promoter;
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Similarity 72.7%;
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99US-0149723P.
99US-0149930P.
99US-0150566P.
99US-0151065P.
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99US-0161350P.
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99US-0161393P.
99US-0161361P.
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                                     Score 40; DB Pred. No. 1.2e 1; Mismatches
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                                DB 3,
1.2e+02;
2;
                                                         Length
                                        Indels
                                                         289;
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RESULT 14
AAG00746
ID AAG00
XX

AAG00746

standard;

protein;

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18-JUN-1999
18-JUN

S-0139456P S-0139458P S-0139458P S-0139461P S-0139461P S-0139462P S-0139763P S-0140339 S-0140339 S-0140353P S-0140353P S-0141083P S-0142950P S-0142950P S-0142950P S-014332P S-0144005P S-0144332P S-0144332P S-0144332P S-0144332P S-0144332P S-0144332P S-0144332P S-0145918P S-0147303P S-0147303P S-0147303P S-0147303P S-0147303P S-0147303P S-014831P S-014831P

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RESULT 15
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                     region (UTR) of the mRNA because they are often obtained from cligo-dr primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' CETs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is a polypeptide encoded by one of a large number of 5. ESTs derived from mRNAs encoding secreted proteins. The 5. ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3. untranslated
                                                         Human; gene
alternative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 13; SEQ ID NO 4827; 71pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and fo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-FEB-2000; 2000EP-00200610.
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     US2003194704-A1
                                                                                                                                  29-JUL-2004
                                                                                                                                                                                        ABO57865 standard; protein; 68
                                                                                                                                                                                                                                                                                                                                                                              Sequence 64 AA;
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                                                                                                     genome
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                                                                                                                                                                                                                                                                                                                                     Similarity
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                                                         expression; single exon probe; microarray; splicing event; genomic alteration.
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                                                                                                   derived single exon protein #4099.
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The invention relates to a nucleic acid probe for measuring human gene expression, comprising any of the 27,400 fully defined nucleotide conciding at least 8 amino acids of any of the 6888 amino acid sequences (fully defined in the specification, or their complements or fragments, and controlled in the specification. The probe is a single exon probe that thybridises under high stringency conditions to a nucleic acid molecule card expression in human cells or tissues. Also included are a spatially-card addressable set of single exon mucleic acid molecule card gene expression (comprising a plurality of single exon nucleic acid gene expression (comprising a plurality of single exon nucleic acid gene expression (comprising a plurality), a single exon microarray for measuring human gene expression, a vector comprising the single exon probe cited above, an ORP-encoded paptide comprising the single exon probe cited above, an ORP-encoded paptide comprising the single exon probe cited above, an ORP-encoded paptide comprising at least 8 contiguous amino acids of any of the above-mentioned amino acid sequences (optionally with conservative amino acid substitutions), an acid customer desiring and/or licensing single exon probes or microarrays to contiguous amino acids of any of the above-mentioned amino acid substitutions), an acustomer desiring and/or licensing single exon probes or microarrays to construct a desiring and/or licensing single exon probes or microarrays to calculate the probes and apparatus are useful in gene capression data by subscription, and a computer-readable cited above. The probes methods and apparatus are useful in gene expression analysis. The probes and apparatus are useful in gene capression of a single exon microarrays. Contains a database having a plurality of records cach record including data on the expression of a single exon microarrays. Contains their probes are useful in dentifying and characterising gross alternative splicing events, in detecting and characterising gross and contains an
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                                                                                                                                                    Sequence 68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 45; SEQ ID NO 31499; 80pp; English
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Search completed: December Job time : 74 secs

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S42098
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capsid protein - cuc
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hypothetical pro
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RESULT 2  UQ1254  Coat protein - cucumber mosaic virus (strain P6)  C;Species: cucumber mosaic virus, CMV  C;Date: 31-Mar-1992 #sequence_revision 05-Aug-1994 #text_ch  C;Accession: JQ1254  R;Shintaku, M.  J. Gen. Virol. 72, 2587-2589, 1991  A;Title: Coat protein gene sequences of two cucumber mosaic  A;Reference number: JQ1253; MUID:92013983; PMID:1919534  A;Recession: JQ1254  A;Molecule type: genomic RNA  A;Molecule type: genomic RNA  A;Residues: 1-218 <shid 1992="" 607-613,="" 92,="" a;rote:="" a;title:="" acetylamino<="" acid="" amino="" amino-terminal="" biochem.="" comparison="" j.="" k.="" micro-identification="" narita,="" of="" prote="" r;tsunasawa,="" s:,="" sequence="" th="" the="" this=""><th>Query Match 100.0%; Score 69; DB 1; Ler Best Local Similarity 100.0%; Pred. No. 3.6e-05; Matches 14; Conservative 0; Mismatches 0; 1  Qy 1 MDKSESTSAGRNRR 14                                     </th><th>RESULT 1 JQ1253 Coat protein - cucumber mosaic virus (strain FC) C;Species: cucumber mosaic virus, CMV C;Date: 31-Mar-1992 #sequence_revision 05-Aug-1994 #text_c; C;Accession: JQ1253 R;Shintaku, M. J. Gen. Virol. 72, 2587-2589, 1991 A;Title: Coat protein gene sequences of two cucumber mosaic A;Reference number: JQ1253; MUID:92013983; PMID:1919534 A;Rccession: JQ1253 A;Molecule type: genomic RNA A;Residues: 1-218 <shi> A;Residues: 1-218 <shi> A;Residues: 1-218 <shi> A;Residues: 1-218 <shi> A;Residues: 1-18; Nuid: RNA A;Cross-references: UNIPROT:Q00259; GB:D10544; NID:g222024 A;Note: comparison of the amino acid sequence of this prot. T,Stunasawa, S.; Narita, K. J. Biochem. 92, 607-613, 1982 A;Title: Micro-identification of amino-terminal acetylamin A;Reference number: A61297; MUID:83056735; PMID:6754709 A;Contents: annotation; acetylation C;Comment: This cucumber mosaic virus strain induces a yel C;Superfamily: cucumber mosaic virus coat protein C;Keywords: acetylated amino end; coat protein C;Keywords: acetylated amino end; Met) #status expe</shi></shi></shi></shi></th><th>ALIGNMENTS</th><th>30 36 52.2 432 2 D86937 31 36 52.2 497 2 C82025 32 36 52.2 671 2 C96546 33 36 52.2 758 2 E81217 34 36 52.2 779 2 T34390 35 36 52.2 1532 2 B81794 36 52.2 1532 2 A86311 37 36 52.2 1908 2 A86311 38 35.5 51.4 1194 1 DJBE28 39 35 50.7 111 2 F27215 40 35 50.7 218 1 B46111 41 35 50.7 218 1 VCVXUV 42 35 50.7 218 1 VCVXUV 43 35 50.7 218 1 VCVXUV 43 35 50.7 218 1 VCVXVIV 44 35 50.7 218 1 VCVXVIV 45 35 50.7 218 1 VCVXVIV 46 35 50.7 218 2 T04758 47 35 50.7 330 2 F72209</th></shid>	Query Match 100.0%; Score 69; DB 1; Ler Best Local Similarity 100.0%; Pred. No. 3.6e-05; Matches 14; Conservative 0; Mismatches 0; 1  Qy 1 MDKSESTSAGRNRR 14	RESULT 1 JQ1253 Coat protein - cucumber mosaic virus (strain FC) C;Species: cucumber mosaic virus, CMV C;Date: 31-Mar-1992 #sequence_revision 05-Aug-1994 #text_c; C;Accession: JQ1253 R;Shintaku, M. J. Gen. Virol. 72, 2587-2589, 1991 A;Title: Coat protein gene sequences of two cucumber mosaic A;Reference number: JQ1253; MUID:92013983; PMID:1919534 A;Rccession: JQ1253 A;Molecule type: genomic RNA A;Residues: 1-218 <shi> A;Residues: 1-218 <shi> A;Residues: 1-218 <shi> A;Residues: 1-218 <shi> A;Residues: 1-18; Nuid: RNA A;Cross-references: UNIPROT:Q00259; GB:D10544; NID:g222024 A;Note: comparison of the amino acid sequence of this prot. T,Stunasawa, S.; Narita, K. J. Biochem. 92, 607-613, 1982 A;Title: Micro-identification of amino-terminal acetylamin A;Reference number: A61297; MUID:83056735; PMID:6754709 A;Contents: annotation; acetylation C;Comment: This cucumber mosaic virus strain induces a yel C;Superfamily: cucumber mosaic virus coat protein C;Keywords: acetylated amino end; coat protein C;Keywords: acetylated amino end; Met) #status expe</shi></shi></shi></shi>	ALIGNMENTS	30 36 52.2 432 2 D86937 31 36 52.2 497 2 C82025 32 36 52.2 671 2 C96546 33 36 52.2 758 2 E81217 34 36 52.2 779 2 T34390 35 36 52.2 1532 2 B81794 36 52.2 1532 2 A86311 37 36 52.2 1908 2 A86311 38 35.5 51.4 1194 1 DJBE28 39 35 50.7 111 2 F27215 40 35 50.7 218 1 B46111 41 35 50.7 218 1 VCVXUV 42 35 50.7 218 1 VCVXUV 43 35 50.7 218 1 VCVXUV 43 35 50.7 218 1 VCVXVIV 44 35 50.7 218 1 VCVXVIV 45 35 50.7 218 1 VCVXVIV 46 35 50.7 218 2 T04758 47 35 50.7 330 2 F72209
_change 09-Jul-2004 aic virus strains reveal a single 26; PIDN:BAA01404.1; PID:g222027 otein to that of strain FC shows ino acids in proteins.	Length 218; Indels 0; Gaps 0;	change 09-Jul-2004 ic virus strains reveal a sin ic virus strains reveal a sin tein to that of strain P6 sho no acids in proteins.		probable membrane probable periplasm probable GTP-bindi organic solvent to hypothetical prote probable solvent t collagen alpha 1 (X protein FIL3.14 (i DNA-directed DNA p BF hand domain prote coat protein - cuc coat protein - cuc coat protein - cuc coat protein - cuc hypothetical prote hypothetical prote hypothetical prote

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A;Reference A;Reference A;Contents: annotation; acceptance virus security C;Comment: This cucumber mosaic virus coat protein C;Superfamily: cucumber mosaic virus coat protein C;Keywords: acetylated amino end; coat protein F;1/Modified site: acetylated amino end (Met) #st: 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              coat protein - cucumber mosaic virus (strain Y) C;Species: cucumber mosaic virus, CMV A;Note: host Nicotiana tabacum cv. Xanthi nc (t C;Date: 30-Jun-1990 #sequence_revision 30-Jun-1 C;Accession: JA0097
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A; Residues: 216-218 < HA2>
A; Residues: 216-218 < HA2>
A; Risunasawa, S;, Narita, K.
J. Biochem. 92, 607-613, 1982
A; Title: Micro-identification of amino-terminal acetylamino acids
A; Reference number: A61297; MUID:83056735; PMID:6754709
A; Contents: annotation; acetylation
C; Genetics:
                                                                                                                                                                                                                                                                                             Ann. Phytopathol. Soc. Jpn. 54,
A;Title: Comparative studies on
A;Reference number: JA0096
A;Accession: JA0097
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C;Superfamily: cucumber mc
C;Keywords: acetylated ami
F;1/Modified site: acetyla
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R;Hayakawa, T.; Mizukai, M.; Nakajima, M.; Suzuki, M.
J. Gen. Virol. 70, 499-504, 1989
A;Title: Complete nucleotide sequence of RNA 3 from cucumber
A;Reference number: JS0089; MUID:89279231; PMID:2732698
                              R;Tsunasawa, S.; Narita, K.
J. Blochem. 92, 607-613, 1982
A;Tille: Micro-identification of amino-terminal acetylamino acids
A;Reference number: A61297; MUID:83056735; PMID:6754709
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A; Residues: 1-218 <NIT>
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A;Residues: 1-218 <HAY>
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Best Local S
Matches 14
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   Contents: annotation;
                                                                                                                                                                                     Cross-references: UNIPROT: P18027
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Accession: JS0090; PS0314
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Pred. No. 3.6e-05;
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A;Gene: cp
C;Superfamily: coa:
C;Keywords: coa:
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A;Title: Coat protein gene sequence analysis of t A;Reference number: JC6073
A;Accession: JC6075
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C;Species: cucumber mosaic virus, CMV
C;Date: 21-Jan-1997 #sequence_revision 21-Jan-1997
C;Accession: JC6075
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A;Aga position: segment RNA3
C;Superfamily: cucumber mosaic virus coat protein
C;Keywords: acetylated amino end; coat protein
C;Keywords: acetylated amino end (Met) #status exp
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                                                                                                                                                                                                                                              submitted to the EMBL Data Library, February 1994 A;Description: Nucleotide sequence of cDNA from R A;Reference number: S42098 A;Accession: S42098
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A; Residues: 1-218 < HYO>
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A; Residues: 1-218 < HUA>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         coat protein
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C.Species: cucumber mosaic virus (strain BS)
C.Species: cucumber mosaic virus, CMV
C.Date: 21-Jan-1997 #sequence_revision 21-Jan-1997
C.Accession: JC6073
R.Huaning
                                                                                                                                                        A;Reference number: JC6073
A;Accession: JC6073
A;Molecula
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S58039
A;Molecule type: mRNA
A;Residues: 1-218 <HUA>
A;Residues: 1-218 <HUA>
A;Cross-references: UNIPROT:Q9DJW8; UNIPROT:Q9YJR7; UNIPROT:Q8JPX1; UNIPROT:Q9E2V6; UNIF
A;Cross-references: UNIPROT:Q9DJW8; UNIPROT:Q66130; UNIPROT:Q8JPX4; UNIPROT:Q9D
EROT:Q91P6; UNIPROT:Q9JFY4; UNIPROT:Q66157; UNIPROT:Q66130; UNIPROT:Q8JPX4; UNIPROT:Q9DW2; UNIPRO
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JC6073
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A;Residues: 1-218 <HAQ>
A;Cross-references: UNIPROT:Q66157; EMBL:X89652; NID:g902619; PIDN:CAA61802.1; PID:g9020
C;Superfamily: cucumber mosaic virus coat protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Gene: cp
C;Superfamily: cucumber mosaic
C;Keywords: coat protein
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Chinese J. Virol. 12, 235-242, 1996
A;Tille: Coat protein gene sequence
A;Reference number: JC6073
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C;Species: cucumber mosaic virus, CMV
C;Date: 21-Jan-1997 #sequence_revision 21-Jan-19
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                                                                                                                                                                                                                                                                              R; Huaping, L.; Hu, J.S.; Barry, K.; Chinese J. Virol. 12, 235-242, 1996
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A; Note: the autic; Genetics:
A; Gene: cp
C; Superfamily: c; C; Keywords: coa
A; Contents:
C; Genetics:
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C;Species: cucumber mosaic virus, CMV
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change
C;Accession: JA0136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Note: this is a revision to the sequence from reference S06932 R;Noel, M.J.T.; Tahar, S.B. Nucletic Acids Res. 17, 10492, 1989 A;Title: Nucleotide sequence of the coat protein gene and flanking, Reference number: S06932; MUID:90098885; PMID:2602158 A;Accession: S06932.
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Nucleic Acids Res. 18, 1332, 1990
A;Title: Corrigendum. Nucleotide sequence of the coat protein A;Reference number: S09663; MUID:90206832; PMID:2320436
                               T:Tsunasawa, S.; Narita, K.
J. Biochem. 92, 607-613, 1982
A;Title: Micro-identification of amino-terminal acetylamino
A;Reference number: A61297; MUID:83056735; PMID:6754709
                                                                                                                                                                        R;Quemada, H.; Kearney, C.; Gonsalves, D.; Slightom, J.L. J. Gen. Virol. 70, 1065-1073, 1989
A;Title: Nucleotide sequences of the coat protein genes a A;Reference number: JU0087; MUID:89279284; PMID:2732712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A, Experimental source: strain I17F
A, Note: this sequence has been revised in
C, Superfamily: cucumber mosaic virus coat
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A;Residues: 1-6,'W',8-44,'W',46-51,'W',53-56,'W',58-68,'W',70,'W',72-73,'W',75-96,'W',98
A;Cross-references: EMBL:X16386
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A; Residues: 1-218 < QUE>
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C;Accession: S09663; S06932
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C; Species: cucumbe A; Variety: strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Experimental source: strain I17F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: UNIPROT:P14767
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Residues: 1-218 < NOE>
                                                                                                     A;Cross-references: UNIPROT: P21368; GB: D00462; NID: g222041;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local
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                  annotation;
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Pred. No. 3.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 69;
Pred. No.
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                                                                                                          PIDN:BAA00357.1;
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A;Cross-references: UNIPROT:Q66120; GB:U20219; NID:g1161916; PIDN:AAC54619.1; PID:g11619A;Experimental source: strain Ixora
C;Comment: The CMV genome consists of three species of single-stranded, capped, positive
C;Comment: The Ixora strain in unusual in that it does not replicate several well charac
ptoms found when replicated by other CMV strains.
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Accession: D71392
R;MCGarvey, P.; Tousignant, M.; Geletka, L.; Cellini, F.; Kaper, J.M.
J. Gen. Virol. 76, 2257-2270, 1995
A;Title: The complete sequence of a cucumber mosaic virus from Ixora
A;Reference number: A71392; MUID:96005047; PMID:7561763
A;Accession: D71392
                                                                                                                                                                                                                                                                                                 R;Barlow, K.
submitted to the EMBL Data Library,
A;Reference number: Z19421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Superfamily: cucumber mosaic virus coat protein C;Keywords: acetylated amino end; coat protein F;1/Modified site: acetylated amino end (Met)/#status predicted
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C;Superfamily: cucumber mosaic virus coat protein
C;Keywords: acetylated amino end; coat protein; glycoprotein
E;1/Modified site: acetylated amino end (Met) #status experimental
F;43/Binding site: carbohydrate (Asn) (covalent) #status predicted
                                                                                    A; Map posi
                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-1711 <WIL>
A; Cross-references: UNIPROT:045409;
A; Experimental source: clone F26H11
                                                                                                                                                                                                                                                        A; Accession: T21432
A; Status: preliminary; translated from
                                                                                                                                                                                                                                                                                                                                                                       hypothetical protein F26H11.2 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 15-Oct.1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004 C;Accession: T21432
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C;Species: cucumber mosaic virus, CMV
C;Date: 03-Aug-1998 #sequence_revision 03-Aug-1998 #text_change 09-Jul-2004
C;Accession: D71392
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A; Residues: 1-218 < MCG>
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                      Query Match
Best Local :
    Matches
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Best Local
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Best Local
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                                                                                      56/3; 154/2; 373/2; 417/3; 505/2; 596/1; 703/2; 847/3;
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13; Conserv
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  Similarity 9; Conserv
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      Conservative
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                    59.4%;
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92.9%;
Score 41; DB:
Pred. No. 42;
1; Mismatches
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Pred. Nd. 0'.0003;
                                                                                                                                                                                               EMBL: Z81515; PIDN: CAB04197.1; GSPDB: GN00020;
                                                                                                                                                                                                                                                                                                                                November 1996
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Search completed: December Job time : 24 secs

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RESULT 14
T31422
C-terminal
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Nucleic Acids Res. 28, 4317-4331, 2000
A;Title: Complete genome sequence of the alkaliphilic bacterium Bacillus halodu: A;Reference number: A83650; MUID:20512582; PMID:11058132
A;Accession: B83765
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B83765
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                                                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-38 <STO>
A; Cross-references: UN
                                                                                                                                                                                                                                                                                                                                                                            hypothetical protein BH0922 [imported] - Bacillus halodurans (strain C-125) C;Species: Bacillus halodurans C;Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 09-Jul-2004 C;Accession: B83765
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R;Yuryev, A.; Patturajan, M.; Litingtung, Y.; Joshi, R.V.; Gentile, Proc. Natl. Acad. Sci. U.S.A. 93, 6975-6980, 1996
A;Title: The C-terminal domain of the largest subunit of RNA polymes A;Reference number: Z21024; MUID:96293459; PMID:8692929
A;Accession: T31422
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change 09-Jul-2004
                                                                                                                                                   A;Gene:
                                                                                                                                                                                     A; Experimental
                                                                                                                                                                                                                                                               A;Status: preliminary
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A; Residues: 1-1473 < YUR>
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                                                                          Matches
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Best Local &
                                                                                                                                                                       Genetics:
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8; Conserv
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                                 MDKSESTSAGRNR 13
MDKKQRMAEGRNR
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                                                                          Conservative
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                                                                                                                                                                                       UNIPROT:Q9KED1; GB:AP001510; GB:BA000004; NID:g10173440; PIDN:BAB0464
Se: strain C-125
                                                                                        55.1%;
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Pred. No.
                                                                                          Score 38;
Pred. No.
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Copyright (c) 1993 - 2004 Compugen Ltd.
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, BB GP4	A.  no L., Kummert J., Belalcazar S., Narvaez J.; no el gen de la proteina de la capside de dos virus del mosaico del pepino (CMV), obtenidos (Musa spp.).";	66133 PRELIMINARY; PRT; 217 AP 66133; 1-NOV-1996 (TrEMBLrel. 01, Created) 1-NOV-1996 (TrEMBLrel. 01, Last sequence v 1-CCT-2003 (TrEMBLrel. 25, Last annotation oat protein. 1-Culmber mosaic virus (cucumber mosaic cucucumber mosaic viruses; seRNA positive-strand viruses, no ucumovirus. 101-1-12305;	y Match 100.0%; Score 69; DB 2; Length 217; Local Similarity 100.0%; Pred. No. 0.00013; hes 14; Conservative 0; Mismatches 0; Indels 0; Gaps 1 MDKSESTSAGRNRR 14	FGTD4;  FRELIMINARY;  PRT;  217 AA.  FGTD4;  5-JUL-2004 (TrEMBLrel. 27, Created)  5-JUL-2004 (TrEMBLrel. 27, Last sequence upda  5-JUL-2004 (TREMBLrel. 27, Last sequence upda  5-JUL-2004 (TREMBLrel. 27, Last annotation up  DAT protein.  CUCUMDER mosaic virus (cucumber mosaic cucumov  iruses; seRNA positive-strand viruses, no DNA  iruses; seRNA positive-strand viruses, no DNA  CBI_TAXID=12305;  1]  CBI_TAXID=12305;  1]  CBI_TAXID=12305;  1]  CBI_TAXID=12305;  1]  CBI_TAXID=12305;  1]  CBI_TAXID=12305;  1]  CBI_TAXID=12305;  CBI_T	

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Matches 14
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R GO; GO:0019028; C:viral capsid; IEA.

R GO; GO:0019028; F:structural molecule activity; IEI

R InterPro; IPR000247; Cucumovirus_coat.

R InterPro; IPR008975; Viral cap_coat.

R Pfam; PR00760; Cucumo coat; 1.

R Pfam; PR00722; CUCUMOCOAT.

R PGDDm; PD001284; Cucumovirus_coat; 1.

Coat protein.
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                         AAR89478;
02-MAR-2004
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01-JUN-2002
01-OCT-2003
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GO; GO:005198; F:structural molecule activity;
InterPro; IPR000247; Cucumovirus coat.
InterPro; IPR008975; Viral_cap_coat.
Pfam; PF00760; Cucumo_coat; 1.
                 Coat
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Submitted (JUL-1995)
EMBL; U32859; AAB501
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ProDom; PD001284; Cucumovirus_coat; 1.
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PIR; JC6075; JC6075.
        Cucumber mosaic virus (cucumber mosaic cucumovirus)
                                                             AAR89478
                                                                                                                                                                              SEQUENCE 217
                                                                                                                                                                                                                                                                            Daolin D., Jie S., Peng Z., Zhixin L., Xiaodong D., Xueqing Z.; "Cloning and Reconstruction of a Expression Vector for the Banana Mosaic Virus Coat Protein Gene.";
                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                Viruses; unclassified viruses.
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Banana mosaic virus
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Pred. No. 0.00013;
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15-DEC-1998
15-DEC-1998
15-DEC-1998
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Hyon S., Park Y.I.;
Submitted (FEB-1994) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: Belongs to the cucumoviruses/bromoviruses coat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Coat protein (Capsid protein) (CP).
Cucumber mosaic virus (strain As) (CMV).
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InterPro; IPR000247; Cucumovirus coat.
InterPro; IPR0008975; Viral_cap_coat.
Pfam; PF00760; Cucumo_coat; 1.
PRINTS; PR00222; CUCUMOCOAT.
PRODOM; PD001284; Cucumovirus_coat; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entities requires a license agreement (Some send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (OCT-2003) to the EMBL/GenBank/DDBJ databases EMBL; AY429437; AAR89478.1; -. Coat protein.
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                                                                1 MDKSESTSAGRNRR 14
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                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                             AA,
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                                                                                                                                                                                                                                                                                24044 MW;
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Pred. No. 0.00013;
                                                                                                                                                                                                                                                                                E6BEC2E3D4CD73EF CRC64;
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                                              This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Suzuki K., Mise K., Inouye N., Okuno T., Furusawa I.;
"Six new subgroup I members of Japanese cucumber mossic virus
determined by nucleotide sequence analysis on RNA3's cDNAs.";
Ann. Phytopathol. Soc. Jpn. 62:40-44(1996).
-!- SIMILARITY: Belongs to the cucumoviruses/bromoviruses coat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Coat protein (Capsid protein) (CP).
Cucumber mosaic virus (strain C7-2) (CMV).
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InterPro; IPR000247; Cucumovirus_coat.
InterPro; IPR008975; Viral_cap_coat.
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"Natural infection of banana by a satellite-containing strain of
cucumber mosaic virus.";
Phytoparasitica 24:49-56(1996).
-!- SIMILARITY: Belongs to the cucumoviruses/bromoviruses coat protein
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Cucumber mosaic virus (strain Banana) (Viruses, ssRNA positive-strand viruses,
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ProDom; PD001284; Cucumovirus_coat; 1.
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(Rel. 37, Last sequence update)
(Rel. 37, Last annotation updat
(Cap. 1, Cap. 1, Ca
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(strain Banana) (CMV).

ve-strand viruses, no DNA stage; Bromoviridae;
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Pred. No.
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RESULT 9
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(Rel. 24, Created)
(Rel. 24, Last sequence update)
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                                                                                                                                                                                                                      the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for cor entitles requires a license agreement (See http://www.isb-sib.ch/aror send an email to license@isb-sib.ch).
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InterPro; IPR008975; Viral_cap_coat.
Pfam; PF00760; Cucumo_coat; 1.
                                                                                                                                      InterPro; IPR000247; Cucumovirus coat.
InterPro; IPR008975; Viral_cap_coat.
Pfam; PF00760; Cucumo_coat; 1.
PRINTS; PR00222; CUCUMOCOAT.
                                                                                                                                                                                                                                                                                                                                           determined by nucleotide sequence analysis on RNA3's cDNAs."; Ann. Phytopathol. Soc. Jpn. 62:40-44(1996).
-i- SIMILARITY: Belongs to the cucumoviruses/bromoviruses coa
                                                                                                                                                                                                                                                                                                                                                                                   Chaumpluk P., Sasaki Y., Nakajima N., Nagano H., Nakamura I.,
Suzuki K., Mise K., Inouye N., Okuno T., Furusawa I.;
"Six new subgroup I members of Japanese cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Coat protein (Capsid protein) (CP).
Cucumber mosaic virus (strain CS) (CMV)
Viruses; ssRNA positive-strand viruses,
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ProDom; PD001284; Cucumovirus_coat;
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                                                                                                    Coat protein.
SEQUENCE 21
                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a obstween the Swiss Institute of Bioinformatics and the EMBL
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                                                                                                                             ProDom; PD001284; Cucumovirus_coat; 1.
                                                                                                                                                                                               EMBL; D28489; BAA05851.1; ~.
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Pred. No. 0.00013;
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15-DEC-1998
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                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                    Chaumpluk P., Sasaki Y., Nakajima N., Nagano H., Nakamura I., Suzuki K., Mise K., Inouye N., Okuno T., Furneawa I.; Suzuki K., et al., Inouye N., Okuno T., Furneawa I.; "Six new subgroup I members of Japanese cucumber mosaic virus determined by nucleotide sequence analysis on RNA3's cDNAs."; Ann. Phytopathol. Soc. Jpn. 62:40-44(1996).
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InterPro; IPR008975; Viral_cap_coat.
Pfam; PF00760; Cucumo_coat; 1.
PRINTS; PR00222; CUCUMOCCAT.
ProDom; PD001284; Cucumovirus_coat;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Coat protein (Capsid protein) (CP).
Cucumber mosaic virus (strain FT) (CMV).
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MEDLINE=92013983; PubMed=1919534;
                                                    HMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cucumovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Viruses; ssRNA positive-strand viruses,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Coat protein (Capsid protein)
Cucumber mosaic virus (strain
                       InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JQ1253; JQ1253.
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(Rel. 37, Last annotation
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                                                 BAA05847.1;
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                    Cucumovirus_coat
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annotation update)
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Pred. No.
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01-APR-1990
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                                    EMBL; X16386; CAA34422.1; -.
EMBL; D10538; BAA01397.1; -.
PIR; S09663; S09663:
InterPro; IPR000247; Cucumovirus coat.
InterPro; IPR0008975; Viral cap coat.
Pfam; PF00760; Cucumo coat; 1.
PRINTS; PR00222; CUCUMOCOAT.
                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMBL outst the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for comentities requires a license agreement (See http://www.isb-sib.ch/an or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                         Owen J., Shintaku M., Aeschleman P., Tahar S., Palukaitis "Nucleotide sequence and evolutionary relationships of cuc virus (CMV) strains: CMV RNA 3.", J. Gen. Virol. 71:2243-2249(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-APR-1990 (Rel. 14, Created)
01-APR-1990 (Rel. 14, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Coat protein (Capsid protein) (CP).
Cucumber mosaic virus (strain I17F) (CMV), an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pfam; PF00760; Cucumo coat; 1.
PRINTS; PR00322; CUCUMOCOAT.
ProDom; PD001284; Cucumovirus_coat;
Coat protein.
                       PRINTS; PR00222; CUCUMOCOAT.
ProDom; PD001284; Cucumovirus_coat; 1.
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                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=90206832; PubMed=2320436; Noel M.J.T., Ben Tahar S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Noel M.J.T., Ben Tahar S.;
"Nucleotide sequence of the coat protein gene cucumber mosaic virus (CMV) strain I17F.";
Nucleic Acids Res. 17:10492-10492(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cucumovirus.
NCBI_TaxID=12308, 12307;
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                                                                                                                                                                                                                                                                                                                                                               STRAIN=FNY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Viruses; ssRNA positive-strand viruses, no DNA stage; Bromoviridae;
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RESULT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Karasawa A., Ito A., Okada I., Hase S., Ehara Y.;

"A possible role of RNA 2 of cucumber mosaic cucumovirus as a
determinant of infection phenotype on cowpea.";
Ann. Phytopathol. Soc. Jpn. 63:289-297(1997).

-!- SIMILARITY: Belongs to the cucumoviruses/bromoviruses coat protein
Paek K.H.;
Submitted (SEP-1994) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: Belongs to the cucumoviruses/bromoviruses
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Cucumber mosaic virus (strain lizuka) (CMV)
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15-DEC-1998 (Rel. 37,
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PRINTS; PR00222; CUCUMOCOAT.
ProDom; PD001284; Cucumovirus_coat; 1.
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InterPro; IPR008975; Viral_cap_coat.
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                                                                                                                                                                                                            InterPro; IPR000247; Cucumovirus_coat.
InterPro; IPR008975; Viral_cap_coat.
Pfam; PF007560; Cucumo coat; 1.
PRINTS; PR00222; CUCUMOCOAT.
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-I- SIMILARITY: Belongs to the cucumoviruses/bromoviruses coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Owen J., Shintaku M., Aeschleman P., 7
"Nucleotide sequence and evolutionary
virus (CMV) strains: CMV RNA 3.";
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InterPro; IPR008975; Viral_cap_coat.
Pfam; PF00760; Cucumo_coat; 1.
                                                                                                                                                                       Coat protein.
SEQUENCE 218
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Cucumber mosaic virus (strain M)
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Q66138;
15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Coat protein (Capsid protein) (CP).
Cucumber mosalc virus (strain N) (CMV).
Viruses; serNA positive-strand viruses, no DNA stage; Bromoviridae;
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InterPro; IRR000247; Cucumovirus_coat.
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Chaumpluk P., Sasaki Y., Nakajima N., Nagano H., Nakamura I.,
Suzuki N., Mise K., Inouye N., Okuno T., Furusawa I.;
Suzuki K., Mise K., Inouye N., Okuno T., Furusawa I.;
"Six new subgroup I members of Japanese cucumber mosaic virus as
determined by nucleotide sequence analysis on RNA3's cDNAs.";
Ann. Phytopathol. Soc. Jpn. 62:40-44(1996).

-I- SINILARITY: Belongs to the cucumoviruses/bromoviruses coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=117123;
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                                                                                                                                                                                                                                               1 MDKSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                  MDKSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       218 AA; 24078 MW; 0BA67C84D8F41B22 CRC64;
                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                        100.0%; Score 69; DB 1; Length 218; 100.0%; Pred. No. 0.00013; tive 0; Mismatches 0; Indels
                                      5, 2004, 08:07:27
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Perfect score:
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                           protein search, using sw model
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length: 2000000000
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
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    GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
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US-08-975-233-10
US-08-975-233-10
US-08-973-352-8049
US-09-328-352-7581
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US-09-328-796A-28057
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Patent No. 542259
Sequence 2, Appli
Sequence 3, Appli
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Sequence 6, Appli
Sequence 15, Appli
Sequence 17, Appli
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Sequence 20, Appli
Sequence 2, Appli
Sequence 4554, Ap
Sequence 19, Appli
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US-08-398-209-2 US-08-398-209-2 US-08-398-209-2 Sequence 2, Application US/08398209 Patent No. 5789656 Patent No. 5789656; Marianne APPLICANT: DeBoth, Michiel APPLICANT: No. 57896566; Marianne APPLICANT: Perret, Joel TITLE OF INVENTION: Transgenic Plants Belonging to the TITLE OF INVENTION: Species Cucumis Melo NUMBER OF SEQUENCES: 4 CORRESPONDENCE ADDRESS: ADDRESSEE: Merchant & Gould STREET: 90 South 7th Street, 3100 No. 5789656west Center CITY: Minneapolis STATE: MN COUNTRY: USA ZIP: 55402 COMPUTER READABLE FORM: MEDIUM TYPE: Flappy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS	RESULT 1 5422259-2 ; Patent No. 5422259 ; PAPLICANT: DE BOTH, MICHIEL; BEN TAHAR, SOPHIA; NOEL, MARIANNE; ; PRITILE OF INVENTION: TRANSGENIC PLANTS BELONGING TO THE ; PATILE OF INVENTION INVEST. ; PAPLICATION NUMBER: US/08/27,563 ; PATILING DATE: 05-MAR-1993 ; PATILING DATE: 13-AUG-1990 ; SEQ ID NO:2: ; PAPLICATION NUMBER: 13-AUG-1990 ; SEQ ID NO:2: ; LENGTH: 217 542259-2  Query Match Best Local Similarity 100.0%; Score 69; DB 6; Length 217; Best Local Similarity 100.0%; Pred. No. 0.00012; Matches 14; Conservative 0; Mismatches 0; Indels 0; G  Qy 1 MDKSESTSAGRNRR 14  Db 1 MDKSESTSAGRNRR 14	28 35.5 51.4 1194 2 US-08-680-326-35 Sequence 3 35.5 50.7 91 4 US-09-252-991A-26033 Sequence 3 30 35 50.7 1163 4 US-09-252-991A-29098 Sequence 3 35 50.7 213 4 US-09-252-991A-30768 Sequence 3 35 50.7 218 3 US-08-875-233-12 Sequence 3 35 50.7 248 4 US-09-270-767-50327 Sequence 3 35 50.7 248 4 US-09-270-767-55327 Sequence 5 50.7 248 4 US-09-270-767-55327 Sequence 5 50.7 266 4 US-09-270-767-55327 Sequence 3 35 50.7 248 4 US-09-270-767-55327 Sequence 5 50.7 248 4 US-09-270-767-55327 Sequence 2 35 50.7 249 4 US-09-270-767-55327 Sequence 3 35 50.7 249 4 US-09-248-796A-19445 Sequence 2 40 35 50.7 499 3 US-09-457-040B-13 Sequence 2 41 35 50.7 571 4 US-09-457-040B-13 Sequence 2 41 35 50.7 571 4 US-09-117-699-6 Sequence 3 50.7 579 4 US-09-117-699-6 Sequence 3 50.7 50.7 50.7 100-08-605-541B-12 Sequence 2 Sequence 3 50.7 50.7 50.9 100-08-605-541B-12 Sequence 2 Sequence 3 Sequence 4 Sequence 3 Sequence 4 Sequence 5 Sequence 6 Sequence
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GENERAL INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/027,563
FILING DATE: 05-MAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/566,255
FILING DATE: 13-AUG-1990
PRIOR APPLICATION NUMBER: FR 8910848
FILING DATE: 11-AUG-1989
ATTORNEY/AGENT INFORMATION:
NAME: Hillson, Randall A.
REGISTRATION NUMBER: 31,838
REFERENCE/DOCKET NUMBER: 8076.21US02
TELECOMMUNICATION INFORMATION:
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TELECOMMUNICATION INFORMATION:
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INFORMATION FOR SEQ ID NO: 2:
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APPLICANT:
APPLICANT:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/027,563
PILING DATE: 05-MAR-1993
PRIOR APPLICATION UNMBER: US 07/566,255
FILING DATE: 13-AUG-1990
                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Perret, Joel
TITLE OF INVENTION: Transgenic Plants Belonging to the
TITLE OF INVENTION: Species Cucumis Melo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOPTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/398,209
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                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
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                                                                                                                                           APPLICATION NUMBER: FILING DATE:
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TYPE: amino acid
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                CITY: Minneapolis
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90 South 7th Street, 3100 No. 5789656west Center
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Ben Tahar, So
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100.0%; Pred. No.
tive 0; Mismatch
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                                                                                                   ; MOLECULE TYPE: protein US-08-553-619B-3
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                                       Best Loc
Matches
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Best Local Similarity
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: DeHaan
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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acids
                                                                                                                                                                                                                                                          FILING DATE: December 1, 199
CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: MARCUS-Wyner, Lynn
REGISTRATION NUMBER: 34,869
REFERENCE/DOCKET NUMBER: 137
                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acids
                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 5919705artis Crop Protection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: DeHaan, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein
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CITY: Palo Alto
STATE: CA
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REGISTRATION NUMBER: 31,838
REFERENCE/DOCKET NUMBER: 8076.21US02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
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TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
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                                                                                                                                                             LENGTH:
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MDKSESTSAGRNRR 14
                                                                                                                                               amino acid
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                                        Conservative
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Virus Resistant Plants
                                                    100.0%;
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ber 1, 1995
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                                                     Score 69; DB 2;
Pred. No. 0.00012;
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MDKSESTSAGRNRR 14

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RESULT 5
US-08-875-233-2
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Patent No.
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Patent No. 6127601
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acids
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/08/875,233
FILING DATE: 26-UN-1997
CLASSIFICATION: 800
CCCANACTON: 800
                                    STREET:
CITY: Chicago
CITY: Illinois
TTATE: IIS.A.
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TITLE OF INVENTION: Plants Resistant to C Strains
TITLE OF INVENTION: Cucumber Mosaic Virus
                                                                                                                                                                                                                                                       APPLICANT: Boeshore, Maury L
APPLICANT: McMaster, J. Russe
APPLICANT: Tricoli, David M
APPLICANT: Reynolds, John F
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TITLE OF INVENTION: Plants Resistant to C Strains
TITLE OF INVENTION: Cucumber Mosaic Virus
                                                                                                                                                 NUMBER OF SEQUENCES: 1
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COMPUTER READABLE FORM:
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STATE: Illinois
COUNTRY: U.S.A.
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GY: linear
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                                                                                                        E: Rockey, Milnamow & Katz, Ltd.
2 Prudential Plaza, 180 N. Stetson, Suite 4700
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McMaster, J. Russell
Tricoli, David M
Reynolds, John F

    Rockey, Milnamow & Katz, Ltd.
    Prudential Plaza, 180 N. Stetson, Suite 4700

                                                                                                                                                                                                                                                                                                Russell
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                                  US-08-875-233-6
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Best Local Similarity
Query Match
                                                                                                                    TELEFAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acids
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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APPLICANT:
APPLICANT:
                                                                                                                                                                       NAME: Lisa V. Mueller TELECOMMUNICATION INFORMATION: TELEPHONE: 312-616-5400
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TITLE OF INVENTION: Plan
TITLE OF INVENTION: Cucu
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                                                  MOLECULE TYPE:
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                                                                   TYPE: ami
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CITY: Chicago
STATE: Illinois
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CLASSIFICATION: 800
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                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/875,233
                                                                                                   LENGTH:
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STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
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amino acid
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3Y: linear
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                                                                                                   218 amino acids
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100.0%;
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Score 69;
 DB 3;
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Length 218;
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Sequence 2, Application US/09127742A
Patent No. 6198022
GENERAL INFORMATION:
APPLICANT: DE BOTH, MICHAEL
APPLICANT: NOEL, MARIANNE
APPLICANT: TAHAR, SOPHIA BEN
APPLICANT: PERRET, JOEL
APPLICANT: BIOSEM
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US-09-127-742-2
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US-08-875-233-15
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TITLE OF INVENTION: TRANSGENIC PLANTS BELONGING
FILE REFERENCE: 8076.21USD2
CURRENT PEPLICATION NUMBER: US/09/127,742A
CURRENT FILING DATE: 1998-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acids
TYPE: amino acid
TOPOLOGY: lina
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APPLICATION NUMBER: US/08/875,233 FILING DATE: 26-JUN-1997 CLASSIFICATION: 800 ATTORNEY/AGENT INFORMATION: NAME: Lisa V. Mueller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Chicago
CITY: Chicago
TTATE: Illinois
TT.S.A.
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICANT: Tricoli, David M
APPLICANT: Reynolds, John F
APPLICANT: Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
TITLE OF INVENTION: Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: 312-616-5400
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CORRESPONDENCE ADDRESS:
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2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                              TO THE SPECIES CUCUMIS
                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 218;
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# EARLIER APPLICATION NUMBER: 08/398,209
# EARLIER FILING DATE: 1995-03-02
# EARLIER APPLICATION NUMBER: 08/027,563
# EARLIER FILING DATE: 1993-03-05
# EARLIER FILING DATE: 1993-03-05
# EARLIER APPLICATION NUMBER: 07/566,255
# EARLIER APPLICATION NUMBER: 07/566,255
# EARLIER APPLICATION NUMBER: 08-13
# EARLIER FILING DATE: 1999-08-11
# NUMBER OF SEQ ID NOS: 4
# SOFTWARE: Patentin Ver. 2.0
# SEQ ID NO 2
# LENGTH: 218
# TYPE: PAT
# ORGANISM: Cucumber mosaic virus
US-09-127-742-2
RESULT 11
$422259 -4
; PATENT NO. 5422259
; APPLICANT: DE BOTH, MICHIEL; BEN TAHAR, SOPHIA; NOEL, MARIANNE;
; PERRET, JOEL
; TITLE OF INVENTION: TRANSGENIC PLANTS BELONGING TO THE
;SPECIES CUCUMIS MELO
; NUMBER OF SEQUENCES: 4
; CURRENT APPLICATION DATA:
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CURRENT FILING DATE: 1998-08-03
EARLIER APPLICATION NUMBER: 08/398,209
EARLIER FILING DATE: 1995-03-02
EARLIER APPLICATION NUMBER: 08/027,563
EARLIER FILING DATE: 1993-03-05
EARLIER APPLICATION NUMBER: 07/566,255
EARLIER APPLICATION NUMBER: 07/566,255
EARLIER FILING DATE: 1990-08-13
EARLIER APPLICATION NUMBER: FR 89 10848
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NUMBER OF SEQ ID NOS: 4
SOFTWARE: PATENTIN VEY: 2.0
SEQ ID NO 4
LENGTH: 218
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: DE BOTH, MICHAEL
APPLICANT: NOEL, MARIANNE
APPLICANT: TAHAR, SOPHIA BEN
APPLICANT: PERRET, JOEL
APPLICANT: BIOSEM
TITLE OF INVENTION: TRANSGENIC PLANTS BELONGING
FILE REFERENCE: 8076.21USD2
                                                                                                                                                                                                                                                                                   1 MDKSESTSAGRNRR 14
                                                                                                                                                                                                                                                      1 MDKSESTSAGRNRR 14
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                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 69; DB 3; 100.0%; Pred. No. 0.00012;
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RESULT 13
US-09-13-999C-4827
; Sequence 4827, Application US/09513999C
; Patent No. 6783961
                                                                                                                                                          Query Match
Best Local Similarity
""" hes 13; Conserv:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              밁
                                                                                                                                                                                                                                            ; LENGTH: 218 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein .
US-08-875-233-10
                                                                                                         용
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                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPARTIBLE
COMPUTER: IBM PC COMPARTIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-UN-1997
CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 10, App-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local
Matches :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: 312-616-5400
TELEFAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/27,563
FILING DATE: 05-MAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 566,255
FILING DATE: 13-AUG-1990
                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Reynolds, John F
APPLICANT: Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
TITLE OF INVENTION: Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 218
                                                                                                                               1 MDKSESTSAGRNRR 14
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: Illinois
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                                                                                                         MDKSESTSAGRNHR 14
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    Rockey, Milnamow & Katz, Ltd.
    Prudential Plaza, 180 N. Stetson, Suite 4700

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Tricoli, David M
                                                                                                                                                                             Conservative
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100.0%; Pred. No. 0.0
tive 0; Mismatches
                                                                                                                                                                           0;
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hes 0;
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Query Match
Best Local Similarity
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US-08-961-564A-5
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US-09-513-999C-4827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961
PILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR PILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SOFTWARE: Patent.pm
SOFTWARE: Patent.pm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 5, Application Patent No. 6114515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/056,935
FILING DATE: 25-AUG-1997
ATTORNEY/AGENT INFORMATION:
NAME: PRESTIA, PAUL F
REGISTRATION NUMBER: 23,031
REFERENCE/DOCKET NUMBER: GH-7023-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-407-0700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: WU, SHUJIAN
APPLICANT: SWEET, RAYMOND
APPLICANT: TRUNEH, ALENGEGED
TITLE OF INVENTION: PIGRL-1, A MEMBER OF IMMUNOGLOBULIN
TITLE OF INVENTION: GENE SUPERFAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 64
TYPE: PRT
                                                                                                    NFORMATION FOR SEQ ID NO:
                                         SEQUENCE CHARACTERISTICS:
LENGTH: 66 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESSE: RATURER & PRESTIA
STREET: P.O. BOX 980
                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/961,564A
FILING DATE: 30-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: P.O. BOX 91
CITY: VALLEY FORGE
                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                               FILING DATE: 30 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 19482
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                                                                                                                                        TELEFAX:
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53 QSESTNNGKNKR 64
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single
linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Diskette
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Pred. No.
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TOPOLOGY: linear
IMMEDIATE SOURCE;
LIBRARY: GenBank
CLONE: 1276649
US-09-382-080-3
                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FRSTSEQ for Windows Version 2.0
CURRENT APPLICATION NUMBER: US/09/382,080
FILING DATE:
PRIOR APPLICATION NUMBER: 08/859,937
FILING DATE:
APPLICATION NUMBER: 08/859,937
FILING DATE:
APPLICATION NUMBER: 36,749
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0308 US
FILECOMMUNICATION INFORMATION:
TELEPHONE: 415-85-0555
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
ILENGTH: 172 amino acids
STRANDEDNESS: single
TOPPOLOGY: linear
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US-09-382-080-3
Search completed: December 5, 2004, 08:07:56 Job time : 24 secs
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Best Local Similarity 72.7
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3, Application US/09382080 Patent No. 6087333
                                                                                                                                     Query Match 55.1%;
Best Local Similarity 70.0%;
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
APPLICANT: Goli, Surya K.
TITLE OF INVENTION: DISEASE ASSOCIATED ACIDIC PROTEIN
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 3174 Porter Drive CITY: Palo Alto COUNTRY. CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                  87 DSTMSGRNRR 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 DKSESTSAGRN 12
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27 DKIETKSAGRN 37
                                                                                                   5 ESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    56.5%; Score 39; DB 3; Length 66; 72.7%; Pred. No. 6.8; tive 1; Mismatches 2; Indels
                                                                                                                                   Score 38; DB 3; Length 172; Pred. No. 29; 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BLOSUM62
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14.144 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1582122 seqs, 356623098 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Published_Applications_AA:*
| /cgn2_6/ptodata/2/pubpaa/USO7_PUBCOMB.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MDKSESTSAGRNRR 14
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Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                 / cgn2_6/ptodata/2/pubpaa/US09E_PUBCOMB.pep:*
/ cgn2_6/ptodata/2/pubpaa/US09E_PUBCOMB.pep:*
/ cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
/ cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
/ cgn2_6/ptodata/2/pubpaa/US10E_PUBCOMB.pep:*
/ cgn2_6/ptodata/2/pubpaa/US10E_PUBCOMB.pep:*
/ cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
/ cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
/ cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
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/cgn2_6/ptodata/2/pubpaa/USO6_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/USO6_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/JSO7_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/USO8_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/USO8_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/USO8_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/USO8_PUBCOMB.pep:*
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39	39	40	40	41	41	41	43	64	69	69	69	69	Score
56.5	56:5	58.0	58.0	59.4	59.4	59.4	62.3	92.8	100.0	100.0	100.0	100.0	Query
103	89	202	159	273	254	193	225	218	218	218	218	218	Query Match Length
17	14	17	15	16	16	16	16	13	13	13	13	13	88
US-10-425-115-188688	US-10-029-386-31499	US-10-425-115-190976	US-10-424-599-184959	US-10-437-963-190100	US-10-437-963-150193	US-10-437-963-135827	US-10-437-963-189118	US-10-011-033-10	US-10-011-033-15	US-10-011-033-6	US-10-011-033-4	US-10-011-033-2	Ω
Sequence 188688,	Sequence 31499, A	Sequence 190976,	Sequence 184959,	Sequence 190100,	Sequence 150193,	Sequence 135827,		Sequence 10, Appl	Sequence 15, Appl	Sequence 6, Appli	Sequence 4, Appli	Sequence 2, Appli	Description

4	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14
37	37	37	37	37	37	37	37	37	37	37	37	37	37	37	37	37	37	38	38	38	38	38	38	38	88	38	38	38	38	38	39
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14	16	17	16	14	15	15	10	9	9	15	16	16	15	16	16	14	16	15	14	15	16	17	15	17	15	14	16	15	17	17	14
US-10-291-265-455	-10-437-963-16986	-10-425-115-3233	0-437-963-13615	-10-369-493-425	-10-374-780A-	0-282-122A-49	US-09-925-302-559	9-925-302-55	-925-300-	-10-424-599-2157	-10-437-963-1107	-10-437-963-1296	-10-425-114-51	-10-767-701-5636	-10-437-	-10-266-886-	-10-4	-10-084-846A	-10-369-493-	-10-424-599-	-10-437-963-19387	-10-425-115-	-10-425-114-	-10-425-115-3	-10-425-114-9	-10-156-761-1107	-10-437-963-10	-10-452-85	25-115-2499	-10-425-115-253	US-10-369-493-4135
Sequence 455, App	Sequence 169865,	32336	13	e 4258,	e 188	e 490	ce 559	e 559, 1	1246	Ø	e 11073	e 12962	(1)	e 56363,	o	æ	æ	6	•	24776	Sequence 193875,	e 20152	e 63801	æ	e 50	e 11072,	e 109979	equence 31,	ው	equence 253470	Sequence 4135, Ap

## ALIGNMENTS

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RESULT 1
US-10-011-033-2
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/10011033 Publication No. US20020124286A1 GENERAL INFORMATION:
                                NAME: Lisa V. Mueller TELECOMMUNICATION INFORMATION: TELEPHONE: 312-616-5400
                                                                                                                                                                                                                                                                                     Reynolds, John r
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Boeshore, Maury L
McMaster, J. Russe
Tricoli, David M
                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                  CITY: Chicago
STATE: Illinois
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                           ADDRESSEE: Rockey, Milnamow & Katz, Ltd. STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                                                           68003
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COMPUTER READABLE FUNCTION
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION NUMBER: US/08/875,233
FILING DATE: 10-000
PRIOR APPLICATION NUMBER: US/08/875,233
FILING DATE: 20-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELEPHONE: 312-616-5400
TELEPHONE: 312-616-5400
TELEPAX: 312-616-5400
TELEPAX: 312-616-5400
TELEPAX: 312-616-5400
TELEPAX: 312-616-5400
TELEPAX: 312-616-5400
TYPE: amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: protein
SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acids
TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
US-10-011-033-4

100.0%; Score 69; DB 13; I
US-10-011-033-4

100.0%; Pred. No. 0.00084;
0: Mismatches 0;
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US-10-011-033-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 MDKSESTSAGRNRR 14
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CITY: Chicago
STATE: Illinois
MDKSESTSAGRNRR 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 218 amino acids
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                                                                                      Length 218;
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US-10-011-033-15
; Sequence 15, Application US/10011033
; Publication No. US20020124286A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 312-616-5400
TELEFAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTIN Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/10/011,033
FILING DATE: 13-NO. US20020124286A1-2001
CLASSIFICATION: <un>
CURNOWN>

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
          CORRESPONDENCE ADDRESS:
ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson,
                                                                                             APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
Carney, Kim J
Carney, Kim J
Cittle Of INVENTION: Plants Resistant to C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
Carney, Kim J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Plants Resistant to C
Cucumber Mosaic Virus
                                                                          NUMBER OF SEQUENCES: 15
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Similarity 100.0%;
14; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 2 Pruder
CITY: Chicago
STATE: Illinois
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CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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Pred. No. 0.00084;
); Mismatches 0;
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                   Suite 4700
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RESULT 5
US-10-011-033-10
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PILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5460
TELEPAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10, Application No. US2003 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local
                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: FISH PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/011,033

FILING DATE: 13-No. US20020124286A1-2001

CLASSIFICATION: CUnknown>

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IENGTH: 218 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compactible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                    APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cucumber Mosaic Virus
NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Reynolds, John F
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
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COMPUTER READABLE FORM:
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                         CITY: Chicago
STATE: Illinois
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                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A.
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100.0%; Pred. No. 0.00084;
tive 0; Mismatches 0;
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RESULT 7
US-10-437-963-135827
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; OTHER INFORMATION: Clone ID: PAT_MRT4530_85657C.1.pep
US-10-437-963-189118
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 189118
LENGTH: 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local (
                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                         Sequence 135827, Application US/10437963 Publication No. US20040123343A1
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 189118, Application US/10437963 Publication No. US20040123343A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                    APPLICANT: La Rosa, Thomas
APPLICANT: Kovalic, David
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: La Rosa, Thomas
APPLICANT: Kovalic, David
APPLICANT: Zhou, Yihua
APPLICANT: Boukharov, Andrey A.
APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                              19 MSNSDKTTAGRHRR 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13;
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                                                                                                                                                                                                                                                                                                                                         1 MDKSESTSAGRNRR 14
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8; Conserv
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                                                                                                 Cao,
Wei
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 218 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 312-616-5400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Boukharov, Andrey A.
Barbazuk, Brad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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David K.
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                                                                                                                                                                                                                                                                                                                                                                                              62.3%;
                                                                                                                                                                                                                                                                                                                                                                                3; Mismatches
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Pred. No. 0.0061;
                                                                                                                                                                                                                                                                                                                                                                                                Score 43;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                DB 16;
                                                                                                                                                                                                                                                                                                                                                                                                                  Length 225;
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CURRENT APPLICATION NUMBER: US/10/437,963

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RESULT 8
US-10-437-963-150193
/ Sequence 150193, Application US/10437963
/ Publication US. US20040123343A1
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US-10-437-963-190100
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US-10-437-963-135827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_50452C.1.pep
US-10-437-963-150193
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Best Local Similarity
""" hes 8; Conserv
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SEQ ID NO 150193
LENGTH: 254
                                                                                                                                                                                                                                                        Sequence 190100, Application US/10437963 Publication No. US20040123343A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 135827
LENGTH: 193
                                                                                                                                                                                                                                           GENERAL INFORMATION:
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APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)8
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
                                                                                                                                                             APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
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ORGANISM: Oryza sativa
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 50.0%;
les 7; Conservation
                                                                                                                                                                                                                                                                                                                                                                    140 LEKSKSTRNGKNKR 153
                                                                                                                                                                                                                                                                                                                                                                                                        1 MDKSESTSAGRNRR 14
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                                                                                                                                        Cao, ...
Wei
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                                                                                                                             Boukharov, Andrey A.
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66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 41; DB Pred. No. 50; 2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 41; DB
Pred. No. 67;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 16; Length 254;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Indels
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Query Match
Best Local Similarity
Farches 9; Conserve
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; OTHER INFORMATION: Clone ID: MRT4577_105749C.1.pep
US-10-425-115-190976
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US-10-424-599-184959
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US-10-437-963-190100
                                                                                            GENERAL INFORMATION:

APPLICANT: La Rosa, Thomas J.

APPLICANT: Kovalic, David K.

APPLICANT: Zhou, Yihua

APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants

FILE REFERENCE: 38-21(5322)B

CURRENT APPLICATION NUMBER: US/10/425,115

CURRENT PILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 369326

SEQ ID NO 190976

LENGTH: 202
                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-425-115-190976
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APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Cao Yongwei
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5)223)B
FILE REFERENCE: 38-21(5)223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 184599
LENGTH: 159
                                                                                                                                                                                                                                                                                                                                                                      Sequence 190976, Application US/10425115
Publication No. US20040214272A1
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Best Local
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LENGTH: 273
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                                                       TYPE: PRT
ORGANISM: Zea mays
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ORGANISM: Glycine max
FEATURE:
                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 40;
Pred. No.
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OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 0.99

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.5

OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 0.89

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.5

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 1.5

OTHER INFORMATION: SWISSPROT HIT: P42285, EVALUE 2.00e-20

US-10-029-386-31499
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                                                               ; OTHER INFORMATION: Clone ID: MRT4577_103673C.1.pep US-10-425-115-188688
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                                                                                                                                                                              APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 188688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
Query Match
Best Local Similarity
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APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: EXPRESSION ANALYSIS TWO FILE REFERENCE: ABOMICA-X-2
CURRENT PILLING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
SOFTWARE: ANDOMAX Sequence Listing Engine vers. 1.1
                                                                                                               LENGTH: 103
TYPE: PRT
ORGANISM: Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                    FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 7; Conserv
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56.5%;
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Score 39;
Pred. No.
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Pred. No. 39;
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  DB 17; Length 103; 59;
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RELICANT: Hinkle, Gregory J.

APPLICANT: Slater, Steven C.

APPLICANT: Goldman, Barry S.

APPLICANT: Goldman, Barry S.

APPLICANT: Chen, Xianfeng

TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF

TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES

FILE REFERENCE: 38-10(52052)B

CURRENT APPLICATION NUMBER: US/10/369,493

CURRENT APPLICATION NUMBER: US/10/369,493

CURRENT APPLICATION NUMBER: US 60/360,039

PRIOR APPLICATION NUMBER: US 60/360,039

PRIOR FILING DATE: 2003-02-21

NUMBER OF SEG ID NOS: 47374

SEQ ID NO 4135

LENGTH: 1104

TYPE: PPT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; ORGANISM: Neurospora crassa US-10-369-493-4135
                                                                                                                                          ; OTHER INFORMATION: Clone ID: MRT4577_162747C.1.pep US-10-425-115-253470
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-425-115-253470
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 15
                                                                                                                                                                                                                           APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE REFERENCE: 38-21(3.222)B
FILE REFERENCE: 38-21(3.222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 253470
LENGTH: 109
                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 253470, Application US/10425115 Publication No. US20040214272A1 GENERAL INFORMATION:
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Best Local
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                                                                                                       Query Match
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ORGANISM: Zea mays
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Pred. No. 6.6e
3; Mismatches
                                                                    Score 38; DB
Pred. No. 93;
5; Mismatches
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Search completed: December

5, 2004, 08:13:54

Job time : 353 secs

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Result
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score
           seq length: 0
seq length: 2000000000
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Match
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(c) 1993 - 2004 Compugen Ltd
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ARX111981
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AR065687
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86.4		86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4	86.4
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A41823 Sequence 4	E01821 cDNA encodi	M22710 Cucumber mo	AJ131623 Cucumber	AJ131624 Cucumber	AR076864 Sequence	A67212 Sequence 1	X77855 Cucumber Mo	U22821 Cucumber mo	X16386 Cucumber mo	BD000279 Transgeni		AR364711 Sequence	AR138343 Sequence	AR021487 Sequence	A39812 Sequence 6	A32133 CMV capsid	BD000278 Transgeni	AR364710 Sequence	AR138342 Sequence	AR021486 Sequence	A32131 CMV capsid	L36525 Cucumber mo	U32858 Cucumber mo	U32859 Cucumber mo	I49963 Sequence 13

## ALIGNMENTS

RESULT 1

Qy	Query Match Best Local : Matches 5	ORIGIN	FEATURES source						COMMENT	OOMMAL	TOTHNAT	TITLE	AUTHORS	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	Locus	BD261798
1 GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACGAGCT 53	Query Match 100.0%; Score 53; DB 6; Length 53; Best Local Similarity 100.0%; Pred. No. 3.6e-11; Matches 53; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	/organism="Cucumber mosaic virus" /mol_type="genomic DNA" /db_xref="taxon:12305"	Location/Qualifiers 1. 53	CC cucumber mosaic virus coating protein peptide rh key Location/Qualifiers FT CDS (6)(47).	Enhancement in protein production by higher plants ubiquitin or	9		PN JP 2002532098-A/2 PD 02-0CT-2002	OS Cucumber mosaic virus	INSTITUTE OF MOLECULAR AGROBIOLOGY	or cucumber mosaic virus coating protein peptide	Enhancement in protein production by higher plants using ubiquitin	Fang, R.X., Wu, J.L. and Chen, X.Y.	1 (bases 1 to 53)	Viruses; ssRNA positive-strand viruses, no DNA stage; Bromoviridae;	Cucumber mosaic virus	Cucumber mosaic virus (cucumber mosaic cucumovirus)	JP 2002532098-A/2.	BD261798:1 GI:33071566	BD261798	Enhancement in protein production by higher plants using ubiquitin or curimber mosaic virus coating protein pentide.		

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REFERENCE
AUTHORS
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AR111975
LOCUS
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JOURNAL
FEATURES
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AR111981
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AR111976
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                                               ARI11975 7
Sequence 3 from patent US
ARI11975
                                                                                                                                                                                       46;
Unknown
                                                                                                                                                                                                                                                                                   Plants resistant to C strains of cucumber mosaic virus Patent: US 6127601-A 14 03-OCT-2000; Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 14 from patent AR111981
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                                     AR111975.1
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Boeshore, M.L., McMaster, R.J., Tricoli, D.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unknown
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Bosehore,M.L., McMaster,R.J., Tricoli,D.M.,
                                                                                                                                                                                                                                                                                                                                                               Unclassified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              h 86.8%; Score 46; DB 6; Lo
Similarity 100.0%; Pred. No. 5.1e-08;
46; Conservative 0; Mismatches 0;
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Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5849548-A 14 15-DEC-1998;
Location/Qualifiers
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 l Similarity 47; Conserv
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M. Cell ablation using trans-splicing ribozymes
Patent: US 5866384-A 14 02-FEB-1999;
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Location/Qualifiers
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1 (bases 1 to 792)
Boesbore, M.L., McMaster, R.J., Tricoli, D.M., Reynolds, J.F.
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nilarity 95.9%;
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                                                            /organism="unknown"
/mol_type="unassigned
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/mol_type="unassigned
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from patent US 5849548.
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M. Cell ablation using trans-splicing ribozymes
Patent: US 5641673-A 14_24-JUN-1997;
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Sequence 14 from patent US
149964
149964.1 GI:2472184
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 6071730-A 14 06-JUN-2000;
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/mol_type="unassigned
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5849548-A 11 15-DEC-1998;
Location/Qualifiers
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M. Cell ablation using trans-splicing ribozymes
Patent: US 6071730-A 11 06-JUN-2000;
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5866384-A 11 02-FEB-1999;
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AR065682
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1 (bases 1 to 152)

Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M. Cell ablation using trans-splicing ribozymes

Patent: US 5866384-A 9 02-FEB-1999;

Location/Qualifiers
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Sequence 11 from patent US 5641673.
149961
149961.1 GI:2472181
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47; Conservat
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Sequence 9 from patent
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AR031570.1 GI:5945859
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M. Cell ablation using trans-splicing ribozymes
Patent: US 5641673-A 11 24-UNN-1997;
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Pred. No. 4.6e-08;
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Sequence 9 from
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Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5849548-A 9 15-DEC-1998;
Location/Qualifiers
                                                                                                                                                                                                                          1 (bases 1 to 152)
Haseloff,J., Brand,A., Perrimon,N.
Cell ablation using trans-splicing
Patent: US 6071730-A 9 06-JUN-2000;
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Aat34666 Coat prot
Aat34665 Coat prot
Aat34665 Coat prot
Aaq10461 Capsid pr
Aaq10462 Capsid pr
Aaq67395 Cauliflow
Aat99545 Cucumber
Aaa795105 Cucumber
Aag76106 Cucumber
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Aat17270 Cucumber
Aaq80683 CMV-SA co
Aaq03641 Cucumber
Aaq80683 CMV-SA co
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Claim 8; Page 18; 42pp; English.

This sequence encodes the C-terminal fragment of a ubiquitin monomer. invention relates to a method for enhancing production of a desired  $\frac{1}{2}$ 

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                                                                                                                                 Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based (known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                    plants, partic. of the family Cucurbitaceae or Sresistant to infection.
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P-PSDB; AAR98895.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACGAGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACGAGCT 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 15 A; 14 C; 13 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first en
                                                                            BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of the V34 strain of cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcmaster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94US-00367789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95WO-US007234.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                         176 A; 199 C; 183 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        . 659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ø
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tricoli DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                  Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 53; DB 3; Pred. No. 3.9e-11; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           V34.
                  46;
                                                                         213 T; 0 U; 0 Other;
                  BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Reynolds JF,
                  2
                                                                                                                                                                                                                                                                                                                                                                                                                                  n DNA - used
Solanaceae,
                  Length 771;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Carney
                                                                                                                                                                                                                                                                                                                                                                                                                                     which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ٥,
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RESULT 4
AAQ10461
ID AAQ1
XX
AC AAQ1
XX

AAQ10461

standard;

976

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RESULT 3
AAT34665
ID AAT3
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                                                          Query Match
Best Local :
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                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
                                                                                                                 Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based c known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                        New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae, resistant to infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-OCT-2003
02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT34665 standard; DNA; 773
                                                                                          Sequence 773 BP; 175 A; 200 C; 185 G; 213 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                   WPI; 1996-333993/33.
                                                                                                                                                                                                                                                                                                                                                        30-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                               07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                     11-JUL-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cucumber mosaic virus; strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT34665;
                                                                                                                                                                                                   Claim 13; Fig 2; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                              WO9621018-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Coat protein;
                                                                                                                                                                                                                                                                                                                                  (ASGR-) ASGROW SEED CO.
                                              1 Similarity
46; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  46;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
              CCATGGACAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 46
CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of the V33 strain of cucumber mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
(first entry)
                                                                                                                                                                                                                                                                                                          Mcmaster JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CP; cucumber mosaic virus; CMV; V27; V33; V34; A35;
                                                                                                                                                                                                                                                                                                                                                        94US-00367789
                                                                                                                                                                                                                                                                                                                                                                               95WO-US007234
                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                        86.8%; Score 46;
100.0%; Pred. No.
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                                              0,
                                                                                                                                                                                                                                                                                                           Tricoli DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   V33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No.
                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.2e-08;
                                                                      DB 2;
                                                          4.2e-08;
                                                                                                                                                                                                                                                                                                          Reynolds JF,
                                                                    Length (773)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            virus
                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                           Carney KJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           49
                                                                                                                                                                                                                                     to produce which are
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                                               Gaps
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RESULT 5
AAQ10462
ID AAQ1
XX
AC AAQ1
XX
AC AAQ1
XY
DT 27-A
DT 25-W
DT 16-A
XX
CMV
XX
CMV
XX
CMV
XX
CMV
XX
CUC\(1000)
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                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-AUG-2003
25-MAR-2003
16-APR-1991
                                                                                                                                                                                                                                                            The gene was isolated from a plasmid (pUC18) containing DNA complementary to RNA 3 of the virulent strain FNY, isolated in New York on infected melons. The sequence includes a leader sequence, coding region and 3 non coding region. The leader sequence is the same length as that of the Japanese Y strain of CMV. Transgenic melon plantlets containing the sequence introduced via an Agrobacterium tumefaciens intermediate are cultured as shoots in special media. The transformed melon plants are resistant to CMV. See also AAQ10462. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PI field.) (Updated on 27-AMR-2003 to correct OS field.)
                     CMV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cbs
                                                                                                                                                                                                                                                                                                                                                                                                  Transgenic melon plantlets prodn. from transformed shoots - culture on specific medium for new transformed plants etc.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-AUG-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EP412912-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CMV; resistance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Capsid
 Cucumber mosaic virus.
                                       Capsid
                                                                   27-AUG-2003
25-MAR-2003
                                                                                                                                                                                                                                            Sequence 976
                                                                                                                                                                                                                                                                                                                                                                     Claim 15; Page 17; 44pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1991-046027/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Deboth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-AUG-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-FEB-1991.
                                                          16-APR-1991
                                                                                               AAQ10462;
                                                                                                                  AAQ10462 standard; DNA; 1007
                                                                                                                                                                                                                                                                                                                                                                                          resistant
                                                                                                                                                                                                                                                                                                                                                                                                                               F-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BIOC-) BIOCEM SA
                    resistance; capsid protein; Cucumis melo;
                                                                                                                                                                  70
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Σ,
                                                                                                                                                                                   Н
                                       protein
                                                                                                                                                                                                      l Similarity
47; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein
                                                                                                                                                                                                                                                                                                                                                                                                                               AAR10652.
                                                                                                                                                                          GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                                                                                                                  GAGTCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bentahar
                                                                                                                                                                                                      86.4%;
llarity 95.9%;
Conservative
                                                                                                                                                                                                                                            B₽;
                                                         (revised)
(revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                          cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene of Cucumber Mosaic Virus strain
                                       gene of Cucumber Mosaic Virus strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  89FR-00010848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90EP-00402282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             capsid protein; Cucumis melo;
                                                                                                                                                                                                                                           224 A; 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             product= "CMV strain FNY capsid"
                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ъ
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Noe1
                                                                                                                                                                                                      Score 45.8; DB 2;
Pred. No. 5.3e-08;
0; Mismatches 2
                                                                                                                                                                                                                                            C; 230
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ξ,
                                                                                                                  ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Perret
                                                                                                                                                                                                                                           <u>ତ୍</u>
                                                                                                                                                                                                                                            273
                                                                                                                                                                                                                                                                                                                                                                                                                                                           4
                                                                                                                                                                                                                                            T; 0 U;
                                                                                                                                                                                                       <u>ب</u>
                     88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88
                                                                                                                                                                                                                       Length 976;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein'
                                                                                                                                                                                                                                           0 Other;
                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FNY
                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                            by two=stage
                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                       0
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RESULT 6
AAQ6730
ID AAQ66
XX AAQ66
XX AAQ66
XX D1 25-M
DT 12-A
XX Caul
XX Caul
XX Caul
XX Caul
XX Caul
XX Caul
XX FF CUS
FT CDS
FT CDS
FT FF CDS
FT T CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The gene was isolated from the virulent French strain 117F of CMV. Tomato plantlets at the 2-leaf stage were infected with CMV strain 117F. 15 days post-infection, the virus was purified from the infected leaves and cDNA was synthesised from RNAs 1,2,3 and 4. After purification and selection by standard techniques, CDNA complementary to RNA 4 was found to encode the capsid protein. It was recloned into "Blue scribe" plasmids and sequenced. Transgenic melon plantlets containing the sequence introduced via an Agrobacterium tumefaciens intermediate are cultured as shoots in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      special media. The transformed melon plants are resistant to CMV. See also AAQ10461. (Updated on 25-MAR-2003 to correct PA field.) (Updated 25-MAR-2003 to correct PA field.) (Updated on 27-AUG-2003 to correct C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Transgenic melon plantlets prodn. from transformed shoots -culture on specific medium for new transformed plants etc., resistant to cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P-PSDB;
                                                                                                                             Cauliflower
                                                                                                                                                                                   Cauliflower mosaic virus;
                                                                                                                                                                                                                         Cauliflower
                                                                                                                                                                                                                                                            25-MAR-2003
12-APR-1995
                                                                                                                                                                                                                                                                                                                    AAQ67395;
                                                                                                                                                                                                                                                                                                                                                       AAQ67395 standard; DNA; 1007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1007 BP; 227 A; 259 C; 247 G; 274 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 16; Page 19; 44pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Deboth M, Bentahar S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-AUG-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-AUG-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-FEB-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EP412912-A.
                                                                                                                                                                   polyribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BIOC-) BIOCEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1991-046027/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
47; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR10653
                                                                                                                                                                                                                                                                                                                                                                                                                                                 GAGTCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                                                                                                               mosaic virus.
                                                                                                                                                                                                                      mosaic virus capsid protein coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                          (revised)
(first entry)
                                                                                                                                                                   inactivate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         89FR-00010848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90EP-00402282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag= a
/product=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 54. .710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                        Location/Qualifiers
                                    /product= "Capsid protein."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      86.4%;
95.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Noel M,
                                                                                                                                                                  CMV; capsid protein; coat protein; inactivation; resistance; crop protection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "CMV strain 117F capsid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         °.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 45.8; D
Pred. No. 5.4e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                         ₽P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Perret
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.
                                                                                                                                                                                                                         sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 1007;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      49
                                                                                                                                                                                                                                                                                                                                                                                                                                                   97
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FR2701960-A1

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RESULT 7
AAT99545/c
ID AAT995
XX AAT995
XX AAT995
XX 25-MAR
DT 21-MAY
XX Coumb
XX FT CDS
FT CDS
FT CDS
FT CDS
FT Promot
FT 
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Best Local S
Matches 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The RNA encoding the capsid protein of cauliflower mosaic virus can be targetted by a nucleic acid sequence called a "polyribozyme". The polyribozyme has endoribonuclease activity and is also to inactivate the gene encoding the viral capsid protein. The polyribozyme comprises several catalytic regions derived from ribozymes and confers complete resistance to virus. See AAQ67391-94. (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New polyribozyme contg. several catalytic regions in complementary sequence - can inactivate gene for viral capsid protein, esp. for proof new virus resistant transgenic plants, also DNA sequence encoding
                                                                                                                                                                                                     promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1994-281767/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lenee P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-FEB-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-SEP-1994.
                   07-MAY-1997;
                                                             12-NOV-1997
                                                                                                        EP806481-A2
                                                                                                                                                                                                                                                                   sig_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Transgenic plant; virus resistance; disease resistance; RNA virus; CMV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cucumber mosaic virus RNA-3 cDNA fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2003
21-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT99545;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT99545 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1007 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-FEB-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GENE-) GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein; antisense gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 47; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Perez P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fig 2; 67pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SHEARS PTY LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
(first entry)
                   97EP-00201379.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93FR-00002269
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                                                                                                                                                                                                                                                                   /product= '956. .1029
                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                   note= "coat
                                                                                                                                                                                                                                             *tag=
                                                                                                                                                                                                                                                                                                                  *tag= b
                                                                                                                                                                                                                                                                                                                                                           *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 227
                                                                                                                                                                                                                                                                                                                                        . 958
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95.9%;
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                                                                                                                                                                                                .1066
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                                                                                                                                                   "F sequence
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                                                                                                                                                                                                                                                                                          "coat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 45.8; DB
Pred. No. 5.4e-(
0; Mismatches
                                                                                                                                                                                                                        protein leader sequence"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88.
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                                                                                                                                                                                                                                                                                          protein'
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                                                                                                                                                   sub-genomic promoter"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5.4e-08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 1007;
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RESULT 8
AAZO7505/c
ID AAZO75
XX AAZO75
XX AAZO75
XX Cucumb
XX Transg
XW CMV; p
XX Cucumb
XX Cucumb
XX Cucumb
XX OB-MAY
XX OB-MAY
XX OB-MAY
XX MPI; 1
XX WPI; 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         have the antisense gene construct integrated into their genomes. They are resistant to viral infection. In particular, they are resistant to CMV. Within the antisense gene construct, the interfering activity of the antisense foomain of the sub-genomic promoter, associated with the antisense activity performed by the coat protein gene, allows production of plants having 100% resistance to CMV. (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This cDNA clone of cucumber mosaic virus (CMV) RNA-3 includes domain F of the subgenomic promoter of viral RNA, the coat protein gene, its leader sequence, and the tRNA-like 3'-terminal region of RNA-3. It was prepared by amplifying a fragment of CMV cDNA-3 from clone pCR-CMVIRNA3 by PCR (see also AAT99548-49). The gene construct is introduced into a vector containing a promoter active in plant cells in antisense orientation relative to the promoter. A claimed recombinant vector comprises the plant promoter, the antisense gene construct and a terminator which is functional in the plant. In addition to CMV, viral RNA may also be used from tobacco mosaic virus and potato virus. Claimed transgenic plants
                                                                                                                                                        09-MAY-1997;
                                                                                                                                                                                                                                                             Cucumber mosaic
                                                                                                                                                                                                                                                                                           Transgenic plant; RNA virus; antisense construct; cucumber mosaic virus; CMV; promoter; coat protein gene; infection; RNA-3; ss.
                                                                                                                                                                                                                                                                                                                                               Cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                 26-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ07505
                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ07505 standard; cDNA; 1067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1066 BP; 299 A; 251 C; 273 G; 243 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Preparing transgenic plants resistant to RNA virus infection - usi: anti:sense gene constructs containing the viral coat protein gene, from cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-538620/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cellini F,
                                                                                                                        09-MAY-1996;
                                                                                                                                                                                          28-SEP-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Page 10; 18pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-MAY-1996;
                                                                                      (META-) METAPONTUM AGROBIOS SCRL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (META-) METAPONTUM AGROBIOS SCRL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      961
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      l Similarity
47; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GAGTCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                        96IT-MI000927
                                                                                                                                                        97US-00854170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95.9%;
                                                                                                                                                                                                                                                                                                                                             (CMV) RNA-3 gene cDNA clone fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 5.4e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 45.8;
Pred. No. 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 1066;
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WPI; 1997-538620/50

Cellini F,

Grieco PD;

V

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RESULT 9
ANNBILLI 1
ID ANNB
AC ANNB
AC ANNB
AC ANNB
AC ANNB
AC ANNB
AC CUCU
XX Synt
KW CUCU
XX Synt
FT CDS

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Best Local S
Matches 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           from the subgenomic promoter, a leader sequence of a coat protein gene of CMV; (c) downstream from the leader sequence, a gene encoding a CMV coat protein; and (d) downstream from the gene, a 3'-terminal region of a CMV coat protein gene. The method is useful for producing plants which are resistant to infection by RNA based viruses. The gene construct gives higher levels of resistance compared to antisense constructs which are capable of complementing with different domains of genomic RNA of CMV. The present sequence represents the fragment of cDNA clone of RNA-3 of CMV. This forms the antisense construct of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to preparing transgenic plants resistant to RNA virus induced infections that comprises integrating an antisense gene construct into the plant genome. The construct comprises (a) an F domai of a subgenomic promoter of cucumber mosaic virus (CMV); (b) downstream
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Preparing transgenic plants resistant to RNA virus infection - using anti:sense gene constructs containing the viral coat protein gene, e.g. from cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                     20-FEB-1987;
25-FEB-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cucumber mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence contg.
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12-NOV-1990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1067 BP; 297 A; 250 C; 276 G; 244 T; 0 U; 0 Other;
                            Disclosure;
                                                                                                                                                                                                                                               Furusawa I,
                                                                                                                                                                                                                                                                                                                                                              18-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-AUG-1988.
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                                                                               producing
                                                                                                                                                                                            1988-236708/34.
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                                                                                                                                                                                                                                                                                                        TAKEDA
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                                                                                                                                                                     AAP80509
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                                                                                                            for
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                            Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; 15pp; English.
                                                                                                                                                                                                                                               Onda H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
                                                                               plants
                                                                                                                                                                                                                                                                                                                                                        87JP-00038288.
87JP-00043443.
88JP-00035809.
                                                                                                         the coat
                                                                                                                                                                                                                                                                                                        CHEM IND LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88EP-00102322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                virus; plant viral resistance; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CMV strain Y coat protein gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /label= cucumber mosaic virus-Y coat protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                         20pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        86.4%;
95.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .1074
                                                                                  resistant
                                                                                                                                                                                                                                               Komiya T;
                                                                                     protein of cucumber mosaic virus strain Y - sistant to cucumber mosaic virus infection.
                            English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 45.8; DB 2;
Pred. No. 5.4e-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                  cucumber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 1067;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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RESULT 10
AAQ76107
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Best Local S
Matches 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This DNA is produced on screening of a plasmid library and is used to transform plant cells which subsequently produce the coat prote ein of cucumber mosaic virus (CMV) strain Y. This produce is not synthesised ir natural plant cells. The resistance to CMV infect- ion, provided by the DNA, is shown in plant cells and redifferent- iated plant bodies. See also AAN81110. (Updated on 25-MAR-2003 to correct PR field.) (Updated on
                                                                                                                    DNA constructs for imparting pathogen, espencedes RNA that interacts with viral RNA
                                                                                                                                                      P-PSDB;
                                                                                                                                                                                                             (SANO)
(SANO)
                                                                                                                                                                                                                                                                                                                                                                        CDS
                                                                                                                                                                                                                                                                                                                                                                                                        Cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                      virus resistance; disease resistance; transgenic plant; cross protection; hypersensitive response; crop improvement; tomato; Lycopersicon esculentum; tobacco; Nicotiana tabacum; RNase T1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cucumber mosaic virus RNA-3 chimera encoding RNase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-MAR-2003
20-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ76107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ76107 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1379 BP; 306 A; 342 C; 333 G; 398
                                                                                                                                                                                        De Haan PT;
                                                                                                                                                                                                                                                          04-JUN-1993;
                                                                                                                                                                                                                                                                                03-JUN-1994;
                                                                                                                                                                                                                                                                                                      22-DEC-1994.
                                                                                                                                                                                                                                                                                                                            WO9429464-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cucumber mosaic virus; tomato mosaic virus; minus-strand RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003 to correct PA field.)
                                                                                                                                                                   1995-036490/05
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                                                                                                                                                                                                                                                                                                                                                                                                                               inhibitory protein;
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                                                                                                                                                                                                             SANDOZ LTD.
SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN
                                                                                                                                                      AAR67754.
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(first entry)
                                                                                                                                                                                                                                                           93GB-00011593
                                                                                                                                                                                                                                                                                94WO-EP001817
                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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/product= "RNase T1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        86.4%;
95.9%;
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Pred. No. 5.7e
O; Mismatches
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                                                                                                                                                                                                                GES
                                                                                                                                                                                                                MBH
                                                                                                                      esp. virus, resistance
NA polymerase to genera
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.7e-08;
les 2;
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A chimeric cucumber mosaic virus RNA-3 (given in AAQ76107) has the operation gene replaced by a gene encoding a cell inhibitory protein, T1, having the sequence given in AAR67754. The construct elicits a resease RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus-resistance on a host plant, e tobacco, tomato. (Updated on 25-MAR-2003 to correct PN field.)

plant, e.g.

a minus-

an

eliciting

agent.

Claim

4; Page 31-32; 50pp;

English.

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RESULT 11

AAQ76106

ID AAQ76

AC COAT

AC CO
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Query Match
Best Local S
Matches 47
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Best Local S
Matches 47
                                                                                                                                                    A chimeric cucumber mosaic virus RNA-3 (given in AAQ76106) codes for the coat protein (CP) of tomato mosaic virus (AAR67752) as well as its own CP (AAR67753). The construct elicits minus-sense RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus-resistance on a host plant, e.g. tobacco, tomato. (Updated on 25-MAR-2003 to correct PN field.)
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(SANO
(SANO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus resistance; disease resistance; transgenic plant; cross protection; hypersensitive response; crop improvement; tomato; Lycopersicon esculentum; tobacco; Nicotiana tabacum; coat protein; CP;
                                                                                                        Sequence 1860 BP; 448 A; 429 C; 431 G; 552 T; 0
                                                                                                                                                                                                                                                                                                                        Claim 3; Page 27-28; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                       DNA constructs for imparting pathogen, esp. virus, resistance to encodes RNA that interacts with viral RNA polymerase to generate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  De Haan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-JUN-1994;
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20-JUL-1995
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                                                                                                                                                                                                                                                                                                                                                                      eliciting
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cucumber mosaic virus RNA-3 chimera encoding ToMV CP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ76106 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            1995-036490/05.
DB; AAR67752, AAR67753.
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              Similarity 47; Conserv
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SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
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                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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(first en
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895. .1550
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  product=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          *tag=
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                                  86.4%;
95.9%;
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95.9%;
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Pred. No. 66
                                  Score 45.8; DB 2;
Pred. No. 6.1e-08;
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0; Mismatches
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                 Mismatches
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                                                        DB 2;
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                                                                                                        U; 0 Other;
                                                        Length 1860;
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RESULT 12
AAQ76108
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                                                                                                     Query Match
Best Local Similarity
Matches 47; Conserv
                                                                                                                                                                                                                                                            A chimeric cucumber mosaic virus RNA-3 (given in AAQ76108) has the coat protein gene replaced by a gene encoding an elicitor, ToMV P30, having the sequence given in AAR67755. The construct elicits a minus-sense RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus-resistance on a host plant, e.g. tobacco, tomato. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 199
P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SANO)
(SANO)
                                                                                                                                                                                                              Sequence 2173 BP; 561 A; 438 C; 532 G; 642 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA constructs for imparting pathogen, encodes RNA that interacts with viral F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           De Haan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-JUN-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus resistance; disease resistance; transgenic plant; cross protection, hypersensitive response; crop improvement; tomato; Lycopersicon esculentum; tobacco; Nicotiana tabacum; P30; elicitor; ToMV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cucumber mosaic virus RNA-3 chimera encoding ToMV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2003
20-JUL-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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DB; AAR67755.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  5; Page 34-35; 50pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SANDOZ LTD.
SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN VERW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PT;
                              GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
GAGTCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 1253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GAGTCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                                                                                        Conservative
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(first en
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/product=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                              86.4%;
95.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    English
                                                                                                     0
                                                                                                     Score 45.8; DB 2;
Pred. No. 6.3e-08;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , esp. virus, resistance to plants RNA polymerase to generate an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           minue-etrand
                                                                                                                                                    Length 2173;
                                                                                                        Indels
                                                    49
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                                                                                                        0
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                                                                                                     Gaps
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RESULT 13 AAT34664

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AAT17259
ID AATT
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AC AATT
XX
AC AATT
DT 16-(
DT 02-1
DT 02-1
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                                                                                                                                                                                                                                                             RESULT 14
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                                                                                                                                                                                                                             Query Match
Best Local S
                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                              Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CNV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based on known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Coat
                                                                             16-OCT-2003
02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                   Claim 2; Fig 1; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                 New isolated cucumber mosaic virus coat protein DNA - used to produce plants, partic. of the family Cucurbitaceae or Solanaceae, which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-OCT-2003
02-DEC-1996
           Cucumber mosaic virus;
                              resistance; ss.
                                        Coat
                                                           Coat protein of the A35 strain of cucumber mosaic virus
                                                                                                           AAT17259;
                                                                                                                               AAT17259 standard; DNA; 772
                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1996-333993/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Boeshore ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-JUL-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9621018-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cucumber mosaic virus; strain V27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           resistance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Coat protein; CP; cucumber mosaic virus; CMV; V27; V33; V34; A35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT34664;
                                                                                                                                                                                                                                                                                                                                                                        resistant to infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ASGR-) ASGROW SEED CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT34664 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein
                                       protein;
                                                                                                                                                                                           4 CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
                                                                                                                                                                                                                      45;
                                                                                                                                                                                                                                                                                                                                                                                                               AAR98893.
                                                                                                                                                                                                                              Similarity
                                                                                                                                                                               CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGGCG
                                                                                                                                                                                                                                                            772 BP; 177 A; 202 C; 182 G; 211 T; 0 U; 0 Other;
                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of the V27 strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
(first entry)
                                       CP;
                                                                             (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcmaster JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9408-00367789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      95WO-US007234.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                       cucumber
                                                                                                                                                                                                                              83.8%;
97.8%;
           strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             772
                                       mosaic virus;
                                                                                                                               BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ₽₽
                                                                                                                                                                                                                     Score 44.4; DB 2;
Pred. No. 1.8e-07;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Tricoli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                           Ŋ,
                                       CMV;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Reynolds JF,
                                       V27;
                                                                                                                                                                                                                                        Length 772;
                                                                                                                                                                                                                     Indels
                                       V33; V34; A35;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Carney KJ;
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RESULT 15
AAN90249
ID AAN90
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Best Local :
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25-MAR-2003
01-NOV-1989
                                                                                                                                                                                                                                                                   AAN90249;
                                                                                                                                                                                                                                                                                           AAN90249 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 772 BP; 175 A; 201 C; 185 G; 211 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of CDNA using primers based cknown CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig 8; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated cucumber mosaic virus coat protein DNA - used to produce plants, partic. of the family Cucurbitaceae or Solanaceae, which are resistant to infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CDS
                         21-DEC-1987;
                                                08-DEC-1988;
                                                                         29-JUN-1989
                                                                                                                           Cucumber mosaic
                                                                                                                                                    solanacea.
                                                                                                                                                               Cucumber mosaic virus C strain;
                                                                                                                                                                                        Cucumber mosaic virus C coat protein gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P-PSDB; AAR93803
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Boeshore ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9621018-A1
                                                                                                  WO8905858-A
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(UPJO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1996-333993/33.
                                                                                                                                                                                                                                                                                                                                                                          4 CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
 UPJOHN
                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                           CCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGGCG
                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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(revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcmaster
                         87US-00135591
                                                  88WO-US004321
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95WO-US007234
                                                                                                                          virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                           DNA; 1423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            . 659
                                                                                                                                                                                                                 entry)
                                                                                                                                                                                                                                                                                                                                                                                                                      83.8%;
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                                                                                                                           strain
                                                                                                                                                                                                                                                                                                                                                                                                         Score 44.4; D
Pred. No. 1.8e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tricoli DM,
                                                                                                                                                                                                                                                                                            ВP
                                                                                                                                                               virus-resistant plants;
                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2;
L.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Reynolds
                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 772;
                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Carney
                                                                                                                                                                cucurbitacea;
                                                                                                                                                                                                                                                                                                                                                           46
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                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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WPI; 1989-206617/28

Quemada H,

Slightom

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AXX
Coat protein gene from C strain of cucumber mosaic virus - used to PT prepare plant transformation vectors and virus-resistant plants.

XX
Claim 1; Page 15; 19pp; English.

XX
Coat protein gene from cucumber mosaic virus C strain is inserted into a CC transformation vector also comprising the 35s promoter of cauliflower cc mosaic virus, and the polyadenylation signal of either the Cauliflower or the Phaseolus vulgaris seed storage gene. This is used to transform cc plants of cucurbitacea and solanacea to produce virus-resistant plants. CC (Updated on 25-MAR-2003 to correct PA field.) (Updated on 24-OCT-2003 to standardise OS field)

XX
Squence 1423 BP; 385 A; 337 C; 320 G; 381 T; 0 U; 0 Other;
Query Match
Best Local Similarity 93.9%; Pred. No. 2.5e-07;
Matches 46; Conservative 0; Mismatches 3; Indels 0; Gaps 0;
Matches 46; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Query Match 370 GAGTCATGGACAAATCTGAATCAACCAGTGGTCGTAACCGTCGACG 49

1 GATCCATGGACAAATCTGAATCAACCAGTGGTCGTAACCATCGACG 49

1 GATCCATGGACAAATCTGAATCAACCAGTGGTCGTAACCATCGACG 49

2 Gaps 0;

Search completed: December 5, 2004, 06:13:16
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Result
No.
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Run on:
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                000
                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          seq length: 0 seq length: 2000000000
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Gapop 10.0 , Gapext 1.0
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Match
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53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     December 5, 2004, 03:35:58; Search time 1476 Seconds (without alignments) 1308.473 Million cell updates/sec
                              EST:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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gb_est2:*
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gb_est6:*
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                              CR153168
CC14018
BX376013
CG880078
CC122836
CD122836
AZ339011
BX23293
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BM067231 KS08001E1
H07817 Khtj009 BN1
AV589910 AV589910
CR153168 Forward 8
CC140418 NDL.2417.
BX376013 BX376013
CG880078 ZMMBBb050
CD122050 ME1-0070P
CD121836 ME1-0070P
AZ339011 IM0070118
BJ232993
BJ36040 387348 MA1
BX922152 BX922152
CN698549 B0409C12-
CC1708124 OR BB8002
CC0703958 DG32-270k
CL733950 OR BB8006
CG802815 ZMMBBb049
CC815885 ZMMBBb049
CC815885 ZMMBBb049
CC815885 ZMMBBb049
CC815885 ZMMBB6052
BM967398 EST30 Tom
BB433897 EST308451
BR435780 EST308451
BR435780 EST308451
BR435780 EST308451
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23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4	23.4
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AW934244	AW442094	BE435313	BE434157	AW221911	BH977843	AW221834	BF113438	BF051011	BE434904	BE436895	BE460690	BE462001	BF112783	AW220906	BE460657	BE436288	BE461053	BE433535	BE434784	TOCOLA
AW934244	AW442094	BE435313	BE434157	AW221911	BH977843	AW221834	BF113438	BF051011	BE434904	BE436895	BE460690	BE462001	BF112783	AW220906	BE460657	BE436288	BE461053	BE433535	BE434784	200001
EST360087	EST311490	EST406391	EST405235	EST298722	ode14c11	EST298645	EST441028	EST436186	EST40598;	EST408013	EST412109	EST413339	EST440373	EST29737	EST412076	EST407366	EST41247	EST400064	EST405862	10.00

## ALIGNMENTS

Qy da	Query Match Best Local Matches 4	RESULT 1 BM067231/C LOCUS DEFINITION ACCESSION VERRION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL COMMENT SOURCE SOURCE FEATURES FEATURES FEATURES FEATURES	
1 GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49	Match 86.4%; Score 45.8; DB 4; Length 127; Local Similarity 95.9%; Pred. No. 3.4e-07; Indels 0; Gaps 0;	BM067231  IGI:22787366  BST.  Capsicum annuum  (bases 1 to 127)  Lee,S., Kim,SY., Chung,YH., Shin,HJ., Goh,SH., Pai,HS.,  Hur,CG. and Choi,D.  Generation of Expressed Sequence Tags from Hot Pepper (Capsicum  annuum L.) and Sequence Analysis in Relation to Hypersensitive  Response Against Pathogen  Unpublished (2001)  Contact: Doll Choi  Genome Research Center and National Center for Genome Information  Korea Research Institute of Bioscience and Biotechnology  P.O. Box 115, Yusong, Taejeon, 305-600, Republic of Korea  Tel: 82-42-860-4340  Tex:	

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RESULT 3
AV589910
LOCUS
                                                                                                                                REFERENCE
AUTHORS
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ACCESSION
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H07817/c
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ORGANISM
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                                                                                                                                                                                                                                                                             AV589910
AV589910 Bos taurus brain fetus
5', mRNA sequence.
AV589910
AV589910.1 GI:9700903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACGAGCT 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                khtj009 BNL1 Brassica
H07817
H07817.1 GI:872639
                                                                      and Sugimoto,Y.

Establishment of a high throughput EST sequencing system using poly(A) tail-removed cDNA libraries and determination of 36,000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: usohn@bh.kyungpook.ac.kr
EST is putatively homologous to
Seq primer: M13 forward.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1995)
Other_ESTs: Arabidopsis dbEST T44633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 200)
Sohn, U., Lee, C.M., Cho, K.H., Jeon, Y.H., Hahn, T.R. and Nam, H.G. CDNAs from Brassica napus (rape)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Brassica.
                                       Nucleic Acids Res.
                                                                                                                              Takasuga, A., Hirotsune, S., Itoh, R., Jitohzono, A.,
                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
                                                                                                                                                                                                                            Bos taurus
                                                                                                                                                                                                                                              Bos taurus (cow)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Laboratory of Molecular Biology
Kyungpook National University
Dept. of Genetic Eng., Kyungpook National Univ., Taegu 702-701, Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Uik Sohn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Brassica napus (rape)
                                                      bovine ESTs
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                                                                                                                                                  (bases 1 to 571)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note=Tvector: pT7T3D; Site_1: NotI; Site_2: EcoRI; POly(A)-mRNA was purified from the leaf of B.napus. CDNA library was constructed from the mRNAs by oligo(dT) priming and directionally cloned from the NotI site in the vector pT7T3D (Pharmacia) to the EcoRI site. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             organism="Brassica napus"
/mol_type="mRNA"
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Pred. No. 28;
0; Mismatches
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cDNA 3', mRNA sequence.
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RESULT 5
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                                                                                                                                         2 ATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCG 46
                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (20-FEB-2004) Sanger Centre, Hinxton, CB10 1SA, UK. http://www.sanger.ac.uk/MICER Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 711)
Adams,D.J., Biggs,P.J., Cox,A.V., Davies,R.M., van der Weyden,L., Jonkers,J., Smith,J., Plumb,R.W., Taylor,R.G., Nishijima,I., Yu,Y.Rogers,J., and Bradley,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CR153168 711 bp DNA linear GSS 06-JUL-2004 Forward strand read from insert in 5'HPRT insertion targeting and chromosome engineering clone MHPN77n06, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Animal Genetics Division
Shirakawa Institute of Animal Genetics
Odakura, Nishigo, Nishi-shirakawa, Fukushima 961-8061, Japan
Tel: 81-248-25-5641
CC140418 807 bp DNA linear GSS 16-APR-2003 NDL.24I7.SP6 Notre Dame Liverpool Aedes aegypti genomic clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CR153168
CR153168.1 GI:49932013
GSS; genome survey sequence; MICER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Single pass sequencing.
This clone was obtained from a polyA-deleted cDNA library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: kazusugi@cocoa.ocn.ne.jp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fax: 81-248-25-5725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Yoshikazu Sugimoto
                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                           /organism="Mus musculus"
/mol type="genomic DNA"
/db_xref="taxon:10090"
/clone="MHPN77006"
/clone_lib="MHPN"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="Bos taurus brain fetus"
/note="Vector: pzL1; Site_1: Sal1; Site_2: Not1; Poly A
was deleted from a Not1 site"
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/mol_type="mRNA"
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lab_host="DH108"
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clone="E1BR009C12"
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73.8%;
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                                                                                                                                                                                              Score 24.2; DB 9;
Pred. No. 1.4e+02;
0; Mismatches 13;
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BP 191 91006 EVRY cedex - France

BP 191 91006 EVRY cedex - France

Email: segref@genoscope.cns.fr, Web: www.genoscope.cns.fr

1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime

1st etrand cDNA was digested with Not I and cloned

into the Not I and EcoR V sites of the pcMVSPORT 6 vector. Library

was normalized. Library was constructed by Life Technologies, a

division of Invitrogen. This sequence belongs to sequence cluster
                                                                                                                                                                                                                                                                                                                                                                                                                   BX376013 Homo sapiens NEUROBLASTOMA COT 25-NORMALIZED Homo CDNA clone CSODCO21YM11 3-PRIME, mRNA sequence.

BX376013
BX376013
GI:46555591
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 32;
                                                                                                                                                                                                 1 (bases 1 to 1061)
Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
Full-length cDNA libraries and normalization
Unpublished (2001)
On May 8, 2003 this sequence version replaced gi:30434740.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Other_GSSs: NDL.24I7.T7
Contact: Brendan Loftus
Department of Eukaryotic Genomics
                                                                                                                                                                                On May 8, 2003 this Contact: Genoscope
                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: enta@tigr.org
Library was provided
Seg primer: SP6
                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Class: BAC ends.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Loftus, B., Shetty, J., Knudson, D. and BAC end sequencing of Aedes aegypti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota;
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CC140418.1 GI:30009473
                                                                                                                                                               Genoscope - Centre National de Sequencage
                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone lib="Notre Dame Liverpool"
/clone="Vector: pBCBAC1; Site 1: Hind III; The library was
prepared from whole body tissue of newly hatched L1 larvae
by David Severson at the University of Notre Dame and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hongbin Zhang"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /strain="Liverpool"
/db_xref="taxon:7159"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Endopterygota; Diptera; Nematocera; Culicoidea; Aedes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              45.7%;
71.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                survey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  by David Severson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 24.2; DB 8;
Pred. No. 1.5e+02;
0; Mismatches 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Diptera; Nematocera;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA"
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CG880078/c
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KEYWORDS
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JOURNAL
                                                           Matches
                                                                          Query Match
Best Local
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CG880078.1
CG880078.1
GSS.
Zea mays
                                                           33;
                                                                                                                                                                                                                                                                                                                                                                                                           190 Frelinghuysen Road, Piscataway,
Tel: 732 445 3801
Fax: 732 445 5735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bharti,A.K., Young,S., Kavchok,S., Keizer,G., Bronzino,A.C. Zohovetz,V., Fuks,G., Yu,Y., Wing,R. and Messing,J. Sequencing of the maize genome at PGIR (2003c) Unpublished (2003) Contact: Bharti,A.K.

Dr. Joachim Messing's lab
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CG880078 1inear 250 bp DNA linear 2MMBBb0504M22f ZMMBBb (HindIII) Zea mays genomic c 2MMBBb0504M22 5', genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                For more information about this cluster, see http://www.genoscope.cns.fr/cdna?s=CSODCO21AGO6NP1&c=2789.f. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD
clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zea mays
                                                                                                                                                                                                                                                                                                                                                    Seq primer: T7
Class: BAC ends
                                                                                                                                                                                                                                                                                                                                                                                          Email: bharti@waksman.rutgers.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2789.f
                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The Plant Genome Initiative at Rutgers, Waksman Institute,
ATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GATCCTTGGCCMMMTGAGGATGATCCATTGCYGGTCYTCATCCAMGA 1055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGA 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (bases 1 to 350)
                                                                                                                                                                                                                                                                                                                                   quality sequence start:
                                                         Conservative
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/clone lib="Homo sapiens NEUROBLASTOMA COT 25-NORMALIZED"
/note="1st strand cDNA was primed with a NotI-oligo(dT)
primer. Five prime end enriched, double-strand cDNA was
digested with Not I and cloned into the Not I and EcoR V
sites of the pCMVSPORT 6 vector. Library was normalized."
                                                                                                                                       /clone lib="ZMMBBb (HindIII;
/note="Vector: pCUGI; Site_1: HindIII;
                                                                                                                                                                                                                                       /organism="Zea mays"
/mol_type="genomic DNA"
/cultivar="B73"
                                                                                                                                                                /clone="ZMMBBb0504M22"
/lab_host="E. coli DH10B"
/clone_lib="ZMMBBb_(HindIII)"
                                                                                                                                                                                                                         db_xref="taxon:4577"
                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /db_xref="taxon:9606"
/clone="CS0DC021YM11"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             organism="Homo sapiens"

mol_type="mRNA"
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                                                                          45.3%;
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                                                       Score 24; DB 9;
Pred. No. 1.5e+02;
0; Mismatches 15
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Pred. No. 1
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..5e+02;
les 13;
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                                                           15;
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                                                                                         Length 350;
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RESULT 9
CD121836
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DEFINITION
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AUTHORS
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CD122050
VERSION
KĘYWORDS
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MEDLINE
PUBMED
                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                 Local
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E 1 (bases 1 to 465)

E Verjovski-Almeida, S., DeMarco, R., Martins, E.A.L., Guimaraes, P.E.M.,
Ojopi, E.P.B., Paquola, A.C.M., Piazza, J.P., Nishiyama, M.Y. Jr.,
Kitajima, J.P., Adamson, R.E., Ashton, P.D., Bonaldo, M.F.,
Coulson, P.S., Dillon, G.P., Farrias, L.P., Gregorio, S.P., Ho, P.L.,
Leite, R.A., Malaquias, L.C.C., Marques, R.C.P., Miyasato, P.A.,
Nascimento, A.L.T.O., Ohlweiler, P.P., Refs, E.M., Ribeiro, M.A.,
Sa, R.G., Stukart, G.C., Soares, M.B., Gargioni, C., Kawano, T.,
Rodrigues, V., Madeira, A.M.B.N., Wilson, R.A., Menck, C.F.M.,
Setubal, J.C., Leite, L.C.C. and Dias-Neto, E.
Transcriptome analysis of the accelomate human parasite Schistosoma
                                                                                                                                                                                        267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       337
         497 bp mRNA linear MEI-0070P-V169-C07-U.B MEI-0070 Schistosoma mansoni MEI-0070P-V169-C07.B, mRNA sequence. CD121836 CD121836.1 GI:34659890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: verjooiq.usp.br
This sequence was derived from the FAPESP Schistosoma mansoni EST
Genome Project. All sequences in the project were assembled and
annotated. This entry and all the assembled sequences can be seen
in the following URL http://bioinfo.iq.usp.br/schisto/
Plate: MEI-0070P-V208 row: 1 column: H.
                                                                                                                                                                                                                                                               33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Dr. Sergio Verjovski-Almeida
Departamento de Bioquimica
Instituto de Quimica - Universidade de Sao Paulo
Av. Prof. Lineu Prestes 748 sala 1200, 05508-900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schistosoma mansoni
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CD122050 465 bp mt ME1-0070P-V208-H01-U.B ME1-0070 SchimE1-0070P-V208-H01.B, mRNA sequence.
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                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                        ACCTATGTATAAATTTGTTTGAACCATAAATGGACATAACCGTCAACG
                                                                                                                                                                                                                           ATCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACG 49
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                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genet.
                                                                                                                                                                                                                                                                                                                                                                          /dev_stage="egg"
/lab_host="Mus musculus"
/clone_lib="ME1-0070"
                                                                                                                                                                                                                                                                                                                                                                                                                                mol_type="mRNA"
/db_xref="taxon:6183"
/clone="MEI-0070P-V208-H01.B"
/sex="mixed_pool"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Schistosoma mansoni"
                                                                                                                                                                                                                                                                                                                                                         note="Vector: pGEM T-easy"
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                                                                                                                                                                                                                                                                               45.3%;
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                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                              Score 24; DB 6;
Pred. No. 1.6e+02;
0; Mismatches 15
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                                                                           cDNA clone
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SOURCE
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AZ339011
LOCUS
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                                                                                                                REFERENCE
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                     TITLE
                                                                                               AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                               267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   33;
                                                                                                                                                                                                           GSS.
                                                                                                                                                                    Mus musculus
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Schistosoma mansoni
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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Verjovski-Almeida, S., DeMarco, R., Martins, E.A.L., Guimaraes, P.E.M., Ojopi, E.P.B., Paquola, A.C.M., Piazza, J.P., Nishiyama, M.Y. Jr., Kitajima, J.P., Adamson, R.E., Ashton, P.D., Bonaldo, M.F., Coulson, P.S., Dillon, G.P., Farias, L.P., Gregorio, S.P., Ho.P.L., Leite, R.A., Malaquias, L.C.C., Marques, R.C.P., Miyasato, P.A., Mascimento, A.L.T.O., Ohlweiler, F.P., Reis, E.M., Ribeiro, M.A., Sa, R.G., Stukart, G.C., Soares, M.B., Gargjoni, C., Kawano, T., Rodrigues, V., Madeira, A.M.B.N., Wilson, R.A., Menck, C.F.M., Setubal, J.C., Leite, L.C.C. and Dias-Neto, E.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleosto; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; I (bases 1 to 751)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Manoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                AZ339011 751 bp DNA linear GSS 29-SEP-200 IMO0770118F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0070L18 F, genomic survey sequence.

AZ339011 751 bp DNA linear GSS 29-SEP-200 DNA library Mus musculus genomic clone UUGCIM0070L18 F, genomic survey sequence.
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This sequence was derived from the FAPESP Schistosoma mansoni EST
Genome Project. All sequences in the project were assembled and
annotated. This entry and all the assembled sequences can be seen
in the following URL http://bioinfo.iq.usp.br/schisto/
Plate: MEI-0070P-V169 row: 7 column: C.
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Departamento de Bioquimica
Instituto de Quimica - Universidade de Sao Paulo
Av. Prof. Lineu Prestes 748 sala 1200, 05508-900 Sao Paulo -
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                                                                                                                                                                                                                                                                                                                 musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /dev_stage="egg"
/lab_host="Mus musculus"
/clone_lib="ME1-0070"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: pGEM T-easy"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Schistosoma mansoni"
/mol_type="mRNA"
/db_xref="taxon:6183"
/clone="ME1-0070P-V169-C07.B"
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0; Mismatches
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                                                                                                                                                                                                                                              Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GSS 29-SEP-2000
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KEYWORDS
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DEFINITION
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BJ232993/c
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                                                                                                                                                                      Triticum aestivum (bread wheat)
Triticum aestivum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30;
                                Expressed genes in Triticum aestivum Unpublished (2002)
                                                                     Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnollophyta; Liliopsida; Poales; Poaceae; Pooldeae; Triticeae; Triticum.

1 (bases 1 to 417)
Cgihara, Y. and Murai, K.
                                                                                                                                                                                                                                                 BJ232993 Y. Ogihara unpublished aestivum cDNA clone whe2o05 5', BJ232993
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High quality sequence stop: 751.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0070 row: L column: 18
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (2000)
                Contact: Tadasu Shin-i
                                                                                                                                                                                                                                   BJ232993.1
                                                                                                                                                                                                                                                                                                                                                                                                              CATAGACTAATCTGAATGAAGCAGAACTTGCCATCACCGT 89
                                                                                                                                                                                                                                                                                                                                                                                                                                              CATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGT 44
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801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gil 4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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For Genetic Resource Information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /lab_host="E. Coli strain XL10-Gold, T1-resistant, /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWD42nv; Purified genomic DNA from nmusculus C57BL/6J (male) was obtained from the JaLaboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Biomedical Polymers Research Bldg.,
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strain="C57BL/6J"
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                                                                                                                                                                                                                                 GI:20050477
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Pred. No. 1.8e+02;
0; Mismatches 10
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PUBMED
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Query Match
Best Local Similarity
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Fahrenkrug,S.C., Smith,T.P.L., Freking,B.A., Cho,J., White,J., Vallet,J., Wise,T., Rohrer,G.A., Pertea,G., Sultana,R., Quackenbush,J. and Keele,J.W.
                                                                                                                                                                                                                                                                     BACKWARD: GTTTTCCCAGTCACGACG plate: 138 row: A column: 5 Seq primer: ATTTAGGTGACACTATAG.
                                                                                                                                                                                                                                                                                                                                                                                                               Single pass sequencing. Bases called and alt trimmed with phred v0.980904.e. Vector identified by cross_match with the -minscore
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Smith TPL USDA, ARS, US Meat Animal Research Center DSDA, 166, Clay Center, NE 68933-0166, USA Tel: 402 762 4366 Fax: 402 762 4390 Email: smith@email.marc.usda.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sus scrofa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sus scrofa (pig)
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387348 MARC 2PIG Sus scrofa
BI360440
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Tel: 81-559-81-6856
Fax: 81-559-81-6855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           National Institute of Genetics
1111 Yata, Mishima, Shizuoka 411-8540, Japan
                                                                                                                                                                                                                                                                                                                                                FORWARD: AGGAAACAGCTATGACCAT
                                                                                                                                                                                                                                                                                                                                                                                           and -minmatch 12 options.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EST cluster assembly
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                                                                                                                                                                                                                                                                                                                                                                       PRimers
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Location/Qualifiers
/organism="Sus scrofa"
/mol type="mRNA"
/mol type="mRNA"
/db_xref="taxon:9823"
/tissue_type="pooled"
/lab_host="MARC 2PIG"
/clone_lib="MARC 2PIG"
/clone_lib="MARC 2PIG"
/note="Vector: pCMV SPORT6; Site_1: Not1; Site_2: Sal1;
Library made from pooled tissue from testis, ovary,
endometrium, hypothalamus, pituitary, and placenta."
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/dev_stage="Feekes' scale 11.2"
/clone_lib="Y. Ogihara unpublished cDNA library, Wh_e"
                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="mRNA"
/cultivar="Chinese Spring'
/db_xref="taxon:4565"
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/mol_type="mRNA"
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Pred. No. 1.9e
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5', mRNA sequence.
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RESULT 14
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DEFINITION
                       FOCUS
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CN698549 631 bp mRNA linear EST 18-MAY-: E0409C12-5 NIA Mouse E11.5 whole embryo cDNA library (Long) Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BX922152 Sus Scrofa library (scan) Sus scrofa cDNA clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Clone distribution: AGENAE Resource centre. Francois PIUMI, Francois.Piumi@jouy.inra.fr, INRA, CEA Radiobiologie et Etude du genome (LREG), Domaine de Vilvert, 78352, Jouy-en-Josas cedex, FRANCE, +33 (0) 1.34.65.28.02, +33 (0) 1.34.65.22.73
Sequence cleaned of vector, adaptator and repetitions. Contact us at sigenasupport@jouy.inra.fr to obtain the chromatogram of this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genetique Animale
Institut National de la Recherche Agronomique
Chemin de Borde-Rouge - Auzeville BP27, 31326 Castanet-Tolosan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bonnet,A., Tosser-Klopp,G., Benne,F., Cabau,C.,
Soares,M., Bonaldo,F. and Hatey,F.
A Pig Normalised Multi-Tissue cDNA Library
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sus scrofa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Plate: 0022 row: e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Tosser-Klopp G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sus scrofa (pig)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BX922152.1 GI:41139000
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                                                                                                                                                             TCCATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACGAGCT 53
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                                                                                                                             TCCATTCAGAAAAGAGAAGCAGCCAGTGGTGGCTGAGACAGTAGAAGAGGT 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nmalia; Eutheria; (bases 1 to 609)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                33 (0) 5.61.28.51.14
33 (0) 5.61.28.53.08
1: tosser@toulouse.inra.fr
                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                       /dev_stage="from embryos to adults"
/clone lib="Sus Scrofa library (scan)"
/clone lib="Sus Scrofa library (scan)"
/note="Ttissues: adipose tissue, brain, kidney, liver,
muscle, ovary, testis, heart, hypothalamus, pancreas,
skin, spleen, thymus, placenta, pituitary gland, seminal
vesicle, small intestine, uterus, adrenals, bulbo uretral
gland, cerebral trunk, pepididymis, female gonad,
gall-bladder, hippocampus, large intestine, male gonad,
melanocytes, stomach, udder"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mol_type="mRNA"
/db_xref="taxon:9823"
/clone="scan0022d.e.22"
/tissue_type="mixed"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Sus scrofa"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                            44.9%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     44.9%;
66.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chordata; Craniata; Vertebrata; Euteleostomi; Cetartiodactyla; Suina; Suidae; Sus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  column: 22.
                                                                                                                                                                                                        Score 23.8; DB 5;
Pred. No. 2e+02;
0; Mismatches 17;
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Pred. No. 2e+02;
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                                                                                                                                                                                                                                                 Length 609;
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                       EST 18-MAY-2004
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TGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGACGAGC 52

Query Match Best Local Matches

Similarity

44.5%;

Conservative

0;

Score 23.6; DB 7; Pred. No. 2.4e+02; 0; Mismatches 14;

Length 631; Indels

0

Gaps

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ACCESSION
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JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sharov,A.A., Piao,Y., Matoba,R., Dudekula,D.B., Qian,Y.,
VanBuren,V., Falco,G., Martin,P.R., Stagg,C.M., Bassey,U.C.,
Wang,Y., Carter,M.G., Hamateni,T., Alba,K., Akutsu,H., Sharova,L
Tanaka,T.S., Kimber,W.L., Yoshikawa,T., Jaradat,S.A., Pantano,S.
Nagaraja,R., Boheler,K.R., Taub,D., Hodes,R.J., Longo,D.L.,
Schlessinger,D., Keller,J., Klotz,B., Kelsoe,G., Umezawa,A.,
Vescovi,A.L., Rossant,J., Kunath,T., Hogan,B.L., Curci,A.,
D'Urso,M., Kelso,J., Hide,W. and Ko,M.S.
Transcriptome analysis of mouse stem cells and early embryos
PLOS Biol. 1 (3), 410-419 (2003)
Contact: Dawood B. Dudekula
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               musculus cDNA clone NIA:E0409C12 IMAGE:30869411 5', mRNA sequence. CN698549 CN698549.1 GI:47467298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           National Institute on Aging/National Institutes of Health 333 Cassell Drive, Suite 4000, Baltimore, MD 21224-6820, USA Email: cdna@lgsun.grc.nia.nih.gov Plate: E0409 row: C column: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1. (bases 1 to 631)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      quality sequence stop: 631
          /clone="NIA:E0409C12 IMAGE:30869411"
/tissue_type="whole embryo including extraembryonic
tissues at 11.5-days postcoitum"
/dev_stage="E11.5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Mus musculus"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'mol_type="mRNA"
'strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lab_host="DH10B"
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354 TGGACAAAGGTGAAGCACACCGTGCTGCACTTAACCCTCACCTGGC 399

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REFERENCE
AUTHORS
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VERSION
KEYWORDS
Search completed: December Job time : 1480 secs
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ORGANISM
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                                                                                                                                                                            Query Match 44.5%;
Best Local Similarity 76.3%;
Matches 29; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OMAP Project
OMAP Project
Unpublished (2004)
Contact: Rod A. Wing
Arizona Genomics Institute
University of Arizona
Forbes Building Room 303, Tucson, AZ 85721-0036, USA
Tel: 520 626 9595
Fax: 520 621 1259
Email: http://genome.arizona.edu
                                                                                                               10 ACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGA 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oryza rufipogon
Oryza rufipogon
Oryza rufipogon
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Eukaryota; Viridiplantae; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
1 (bases 1 to 666)
1 (bases 1 to 666)
1 Kim,H., Yu,Y., Stum,D., Yost,D., Rao,K., Luo,M., Jetty,R.,
Kudrna,D., Muller,C., Hatfield,J., Soderlund,C. and Wing,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CL708124
666 bp DNA linear GSS 26-JUL-2004
OR BBa0078C18.r OR BBa Oryza rufipogon genomic clone OR BBa0028C18
3', genomic survey sequence.
CL708124
CL708124.1 GI:50595162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR PRIMET'S TAA TAC GAC TCA CTA TAG GG FORWARD: TAA TAC GAC TCA TTA GGC ACC CCA INSERT Length: 161 Std Error: 0.00 Plate: 0028 row: C column: 18 Seq primer: CAC TCA TTA GGC ACC CCA Class: BAC ends.
                                                                                        AMAMATTTTAATCCACAACTGCTGGTCGTGACCGATGA 496
                                                                                                                                                                                                                                                                              /organism="Oryza rufipogon"
/mol type="genomic DNA"
/mol type="genomic DNA"
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/clone="OR_BBa0028C18"
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/lab host="DH108-T1 phage resistant"
/clone_lib="OR_BBa"
/note="Vector: pAGIBAC1; Site_1: HindIII; Site_2: HindIII"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ocation/Qualifiers
                    5, 2004, 06:38:00
                                                                                                                                                                          Score 23.6; DB 9; Length 6 Pred. No. 2.5e+02; O; Mismatches 9; Indels
                                                                                                                                                                                                                 Length 666;
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-MODEL-frame+ p2n.model -DEV=xlh
-Q=/Cgn2 1/USPTO_spool/US09857881/runat_03122004_160551_18585/app_query.fasta_1.199
-Q=/Cgn2 1/USPTO_spool/US09857881/runat_03122004_160551_18585/app_query.fasta_1.199
-DB=GenEmbl -QFWT=fastap -SUFFIX=p2n.rge -MINMATGH=0.1 -LOOPEXT=0
-UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
-DOCALIGN=200 -THR_SCORE=pct -THR MAX=100 -THR MIN=0 -ALIGN=15 -MODE=LOCAL
-OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER=US09857841_@CGN_1 1_3731_@runat_03122004_160551_18585 -NCPU=6 -ICPU=3
-NO_MMAD -LARGEOÜERY -NEG SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                       Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence:
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                                                                                                   No.
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                                                                                                 Score
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
      69 69
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1 MDKSESTSAGRNRR
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### ALIGNMENTS

						COMMENT		JOURNAL		TITLE	AUTHORS	REFERENCE			ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	Locus	BD261798	RESULT 1
CC Enhancement in protein production by higher plants using CC ubiquitin or CC cucumber mosaic virus coating protein peptide FH Key	212	PI RONG XIANG FANG, JUNG LIN WU, XIAO YING CHEN PC C12N15/09.A01H5/00.C07K14/415.C07K19/00.C12N5/10.C12N15/00. PC	Ī	_	PN JP 2002532098-A/2	OS Cucumber mosaic virus	INSTITUTE OF MOLECULAR AGROBIOLOGY	Patent: JP 2002532098-A 2 02-OCT-2002;	or cucumber mosaic virus coating protein peptide	Enhancement in protein production by higher plants using ubiquitin	Fang, R.X., Wu, J.L. and Chen, X.Y.	1 (bases 1 to 53)	Cucumovirus.	Viruses; ssRNA positive-strand viruses, no DNA stage; Bromoviridae;	Cucumber mosaic virus	Cucumber mosaic virus (cucumber mosaic cucumovirus)	JP 2002532098-A/2.	BD261798.1 GI:33071566		or cucumber mosaic virus coating protein peptide.	Enhancement in protein production by higher plants using ubiquitin	BD261798 53 bp DNA linear PAT 17-JUL-2003		

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        1 (bases 1 to 130)
HaselOff,J., Brand,A., Perrimon,N. and Goodman,H.
Cell ablation using trans-splicing ribozymes
Patent: US 5849548-A 14 15-DEC-1998;
Location/Qualifiers
                                                                                                             AR065687
Sequence 14 from patent
AR065687
AR065687.1 GI:5995903
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HaselOff,J., Brand,A., Perrimon,N. and Goodman,H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5866384-A 14 02-FBB-1999;
Location/Qualifiers
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RESULT 4
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Sequence
AR097446
                                                            1 (bases 1 to 130)
Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M. Cell ablation using trans-splicing ribozymes
Patent: US 5641673-A 14 24-UN-1997;
Location/Qualifiers
                                                                                                                                                                                                                                      1 (bases 1 to 130)
Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M. Cell ablation using trans-splicing ribozymes
Patent: US 6071730-A 14 06-JUN-2000;
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5866384-A 11 02-FBB-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unknown.
                                                                                                            1 (bases 1 to 131)
Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5849548-A 11 15-DEC-1998;
Location/Qualifiers
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                                                                                                                                                                                            1 (bases 1 to 131)
Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M. Cell ablation using trans-splicing ribozymes
Patent: US 5641673-A 11 24-JUN-1997;
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RESULT 12
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AR031570
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Haseloff, J., Brand, A., Perrimon, N. al
Haseloff, J., Brand, A., Perrimon, N. al
Cell ablation using trans-splicing r.
Patent: US 5866384-A 10 02-FEB-1999;
Location/Qualifiers
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Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5866384-A 9 02-FBB-1999;
Location/Qualifiers
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1 (bases 1 to 152)

1 (bases 1 to 152)

1 Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M.

Cell ablation using trans-splicing ribozymes

Patent: US 5849548-A 9 15-DEC-1998;

Location/Qualifiers
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M. Cell ablation using trans-splicing ribozymes
Patent: US 5849548-A 10 15-DEC-1998;
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                                         1 (bases 1 to 152)
1 (bases 1 to 152)
Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M. Cell ablation using trans-splicing ribozymes
Patent: US 6071730-A 10 06-JUN-2000;
1 120
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Sequence 10 from patent US 6071730.
AR097442 GI:12806172
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1 (bases 1 to 152)

1 (bases 1 to 152)

Haseloff, J. Brand, A., Perrimon, N. and Goodman, H.M.

Cell ablation using trans-splicing ribozymes

Patent: US 6071730-A 9 06-JUN-2000;

Patent: US 6071730-A 9 16-JUN-2000;

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-DEV TIMEOUT=120 -WARN TIMEOUT=30 -THRADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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AAT99545
AAZ07505
                           AAQ48414
AAT07648
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AAN90249
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 AAT00644
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ABZ41931
ADO61972
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Aat72272 Cucumber
Ab158209 Agrobacte
Ab237724 N. gonorr
Ab237724 N. gonorr
Ab661972 Transcrip
Aca38515 Prokaryot
Abd15266 Pseudomon
Aca40580 Prokaryot
Abd15266 Pseudomon
Aca40580 Prokaryot
Abd15266 Pseudomon
Continuation (23 o
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Aaf66292 Novel hum
Ada72516 Rice gene
Ada72517 Human ost
Acc50398 Arabidops
Asg41742 Cucumber
Aag80683 CMV-SA co
Aac34260 Arabidops
Aad17271 Human inf
Abd17277 Human ade
Abz97741 Human ade
Abz97741 Human ade
Abz97741 Human ade
Abz97741 Human ade
Aat29299 Human ven
Aat00644 Human A1
Aat07648 Human A1
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Aaq76108 Cucumber
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Aaq76107 Cucumber
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## ALIGNMENTS

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RESULT 1
AAA30858
AAA30858 standard; DNA;
                              11-DEC-1998;
                                                                                                                            Nicotiana tabacum.
                                                                                                                                                                  Ubiquitin monomer C-terminal fragment coding sequence.
                                                                                                                                                                                  19-SEP-2000
                                                                                                                                                                                                  AAA30858;
(MOLE-) INST MOLECULAR AGROBIOLOGY.
               11-DEC-1998;
                                              22-JUN-2000
                                                                                                                                                   Obiquitin
                                                             WO200036129-A1
                                                                                                                                                   monomer;
                                                                                                                                                                                  (first entry)
               98WO-SG000103
                              98WO-SG000103
                                                                            /*tag= a
/product=
/partial
                                                                                                            Location/Qualifiers
                                                                                                                                                   protein production; plant cell; ubiquitin promoter;
                                                                                                                                                                                                                  53
                                                                                   "Ubiquitin_monomer_fragment"
                                                                                                                                                                                                                  BP.
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RESULT 2
AAT34666
Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence encodes the C-terminal fragment of a ubiquitin monomer. The invention relates to a method for enhancing production of a desired protein in a plant or plant cell by inserting a nucleic acid (NA) encoding a ubiquitin monomer upstream of a NA encoding the desired protein, where the fusion construct encodes a fusion protein and expression is not controlled by the ubiquitin promoter. The invention also relates to a NA acid vector a NA vector able to transform a plant cell, that comprises NA encoding a fusion protein having a ubiquitin monomer linked to a protein of interest and further, where expression of the fusion construct is not under control of a ubiquitin promoter. The construct allows enhanced production of the desired protein in plants or
                                                                                                                                                                                                                                   Coat protein; CP; resistance; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fang
                                                                                       07-JUN-1995;
                                                                                                                 11-JUL-1996
                                                                                                                                       WO9621018-A1
                                                                                                                                                                                                            Cucumber mosaic virus; strain V34.
                                                                                                                                                                                                                                                                         Coat protein of the V34 strain of cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                 16-OCT-2003
02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Production of desired protein in plants or plant cells ubiquitin monomer coding sequence upstream of the gene
                 Boeshore ML,
                                                                                                                                                                                                                                                                                                                                                          AAT34666 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 53 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    desired protein.
                                         (ASGR-) ASGROW SEED CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000-431604/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8; Page 18; 42pp; English.
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(first entry)
                 Mcmaster JR,
                                                                9408-00367789
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Matches:
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Indels:
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                 Reynolds
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RESULT 3
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        Claim
                                           New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae,
                                                                                  WPI; 1996-333993/33.
P-PSDB; AAR98893.
                                                                                                                                                                                                                                                                                                                                                                                                          16-OCT-2003
02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT34664;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1996-333993/33.
P-PSDB; AAR98895.
                                                                                                                                                                       30-DEC-1994;
                                                                                                                                                                                               07-JUN-1995;
                                                                                                                                                                                                                       11-JUL-1996.
                                                                                                                                                                                                                                                                                                                       Cucumber mosaic virus; strain V27.
                                                                                                                                                                                                                                                                                                                                                           Coat protein; CP; cucumber mosaic virus; CMV; V27; V33; V34; A35;
                                                                                                                                                                                                                                                                                                                                                                                  Coat protein of the V27 strain of cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT34664 standard; DNA;
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                                  resistant to infection.
                                                                                                                      Boeshore ML,
                                                                                                                                              (ASGR-) ASGROW
                                                                                                                                                                                                                                                WO9621018-A1
                                                                                                                                                                                                                                                                                                                                               resistance;
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          <u>ب</u>
       Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
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(first en
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                                                                                                                      Mcmaster JR,
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        80pp; English
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Score:
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XX WO99
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Query Match:
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                                                                                                                                                                                                                                                                                                                         New isolated cucumber mosaic virus coat protein DNA - used to produce plants, partic. of the family Cucurbitaceae or Solanaceae, which are resistant to infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 772 BP; 177 A; 202 C; 182 G; 211 T; 0 U; 0 Other;
                                                         Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based oknown CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-JUN-1995;
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02-DEC-1996
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   Sequence
                                                                                                                                                                                                                                                             Disclosure; Fig 8; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Boeshore ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-DEC-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cucumber mosaic virus; strain A35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Coat protein of the A35 strain of cucumber mosaic virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein; CP; cucumber mosaic virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                           1996-333993/33.
DB; AAR93803.
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      772
      B₽;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mcmaster JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEED CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94US-00367789.
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   175 A;
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      201 C;
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      185 G;
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Matches:
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      211
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      T; 0
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RESULT 5
AAT34665
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        Best Local Similarity:
Query Match:
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Query Match:
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                                                         Alignment Scores:
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                                                                                               Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based c known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae,
                                                                                                                                                                                                                                                                                                                                                                                      CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Coat
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02-DEC-1996
                                                                            Sequence
                                                                                                                                                                  Claim 13; Fig 2; 80pp; English.
                                                                                                                                                                                                                                      WPI; 1996-333993/33.
                                                                                                                                                                                                                                                                                                30-DEC-1994;
                                                                                                                                                                                                                                                                                                                    07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                      11-JUL-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                   Cucumber mosaic virus; strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT34665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT34665 standard; DNA; 773
                                                                                                                                                                                       resistant to infection.
                                                                                                                                                                                                                                                        Boeshore ML, Mcmaster JR,
                                                                                                                                                                                                                                                                                                                                                          WO9621018-A1
                                                                                                                                                                                                                                                                             (ASGR-) ASGROW SEED
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                                                                            BP; 175 A; 200
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                CP; cucumber mosaic virus; CMV; V27; V33; V34;
                                                                                                                                                                                                                                                                                                94US-00367789
                                                                                                                                                                                                                                                                                                                    95WO-US007234
                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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                                                                             185
                                     Length:
Matches:
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Indels:
                  Conservative:
Mismatches:
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          Indels:
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US-09-857-841-4 (1-14) x AAT34665 (1-773)

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RESULT 6
AAQ10461
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                                                   US-09-857-841-4 (1-14) x AAQ10461 (1-976)
                                                                         Query Match:
DB:
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Best Local Similarity:
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                                                                                                                                                                                     The gene was isolated from a plasmid (pUC18) containing DNA complementary to RNA 3 of the virulent strain FNY, isolated in New York on infected melons. The sequence includes a leader sequence, coding region and 3 non-coding region. The leader sequence is the same length as that of the Japanese Y strain of CMV. Transgenic melon plantlets containing the sequence introduced via an Agrobacterium tumefaciens intermediate are cultured as shoots in special media. The transformed melon plants are resistant to CMV. See also AAQ10462. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PI field.) (Updated on 27-AMR-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                Claim
                                                                                                                                                                                                                                                                                                                                               Transgenic melon plantlets prodn. from transformed shoots - culture on specific medium for new transformed plants etc.,
                                                                                                                                                                                                                                                                                                                                                                                                                     Deboth M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Capsid protein gene of Cucumber Mosaic Virus strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-AUG-2003
25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ10461;
                                                                                                                                                                                                                                                                                                                                        resistant to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EP412912-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            (BIOC-) BIOCEM SA
                                                                                                                                                                                                                                                                                                                                                                                               1991-046027/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      resistance; capsid protein; Cucumis melo;
                                                                                                                                                                                                                                                                                                               15; Page 17; 44pp; French.
       75
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                 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
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                                                                                                                                                                                                                                                                                                                                                                                                                     Bentahar S,
                                                                                                                                                                  BP;
                                                                                                                                                                                                                                                                                                                                       cucumber mosaic virus.
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                                                                                                                                                                 224 A; 249
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100.00%
100.00%
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Matches:
Conservative:
Mismatches:
Indels:
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14
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     116
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RESULT 7
AAQ10462
RESULT 8
                                                                                                                      US-09-857-841-4 (1-14)
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Best Local Similarity:
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                                                                                                                                                                                                                                                 Score:
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25-MAR-2003
16-APR-1991
                                                                                                                                                                                                                                                                                                                                                                                 The gene was isolated from the virulent French strain 117F of CMV. Tomato plantlets at the 2-leaf stage were infected with CMV strain 117F. 15 days post-infection, the virus was purified from the infected leaves and cDNA was synthesised from RNAs 1,2,3 and 4. After purification and selection by standard techniques, cDNA complementary to RNA 4 was found to encode the capsid protein. It was recloned into "Blue scribe" plasmids and sequenced. Transgenic melon plantlets containing the sequence introduced via an Agrobacterium tumefaciens intermediate are cultured as shoots in special media. The transformed melon plants are resistant to CMV. See also AAQ10461. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct OS
                                                                                                                                                                                                                                                                                                                              Sequence 1007 BP; 227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Transgenic melon plantlets prodn. from transformed shoots - by t culture on specific medium for new transformed plants etc., esp. resistant to cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1991-046027/07.
P-PSDB; AAR10653.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Deboth M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-AUG-1989;
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                                      54
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                                                             MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
                                      ATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGA 95
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(revised)
(first en
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/product= "CMV strain l17F capsid protein"
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                                                                                                                    (1-1007)
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Matches:
Conservative:
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                                                                                                                                                                                  Mismatches:
Indels:
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RESULT 9
AAT99545/c
ID AAT995
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AC AAT995
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AC AAT995
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DT 25-MAR
DT 21-MAY
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The RNA encoding the capsid protein of cauliflower mosaic virus can be targetted by a nucleic acid sequence called a "polyribozyme". The polyribozyme has endoribonuclease activity and is able to inactivate the gene encoding the viral capsid protein. The polyribozyme comprises several catalytic regions derived from ribozymes and confers complete resistance to virus. See AAQ67391-94. (Updated on 25-MAR-2003 to correct by field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New polyribozyme contg. several catalytic regions in complementary sequence - can inactivate gene for viral capsid protein, esp. for proof new virus resistant transgenic plants, also DNA sequence encoding
  25-MAR-2003
21-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Fig 2; 67pp;
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12-APR-1995
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                                                                           AAT99545;
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                                                                                                                        standard; cDNA; 1066
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e; inactivate;
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crop protection;
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                                                                                                                                                                     09-MAY-1996;
                                                                                                                                                                                          12-NOV-1997
                                                                                                                                                                                                    EP806481-A2
                                                                                                                                                                                                                                                                                                            Cucumber mosaic virus RNA-3 cDNA fragment
                                                                                                                                                                                                                                        sig_peptide
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                                                                                                                                                           (META-) METAPONTUM AGROBIOS SCRL
                                                                                                                                                                                                                                                                                            sgenic plant; virus protein; antisense
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                                                                                                                                                                               97EP-00201379
                                                                                                                                                                     96IT-MI000927.
                                                                                                                                                                                                                                                                                   virus
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300.
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                                                                                                                                                                                                               note= "F sequence of sub-genomic promoter"
                                                                                                                                                                                                                              note= "coat
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                                                                                                                                                                                                                                                         . 958
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                                                                                                                                                                                                                                                                                                  resistance; disease resistance; RNA virus; CMV;
                                                                                                                                                                                                                                                                                             gene;
                                                                                                                                                                                                                                              "coat
                                                                                                                                                                                                                              protein leader sequence
                                                                                                                                                                                                                                              protein"
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Preparing transgenic plants resistant to RNA virus infection - using anti:sense gene constructs containing the viral coat protein gene, e. cucumber mosaic virus. Ġ

Claim 3; Page 10; 18pp; English.

This cDNA clone of cucumber mosaic virus (CMV) RNA-3 includes domain F of the subgenomic promoter of viral RNA, the coat protein gene, its leader sequence, and the tRNA-1ike 3'-terminal region of RNA-3. It was prepared by amplifying a fragment of CMV cDNA-3 from clone pCR-CMVIRNA3 by PCR (see also AAT99548-49). The gene construct is introduced into a vector containing a promoter active in plant cells in antisense orientation relative to the promoter. A claimed recombinant vector comprises the plant promoter, the antisense gene construct and a terminator which is functional in the plant. In addition to CMV, viral RNA may also be used from tobacco mosaic virus and potato virus. Claimed transgenic plants have the antisense gene construct integrated into their genomes. They are resistant to viral infection. In particular, they are resistant to CMV. Within the antisense gene construct, the interfering activity of the antisense F domain of the sub-genomic promoter, associated with the antisense activity performed by the coat protein gene, allows product of plants having 100% resistance to CMV. (Updated on 25-MAR-2003 to correct PR field.) production F of are

Sequence 1066 BP; 299 P. 251 <u>ر</u> 273 <u>ი</u> 243 Η. 0 U; 0 Other;

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Percent Similarity:
Best Local Similarity:
Query Match:
US-09-857-841-4 (1-14) x AAT99545 (1-1066)
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Matches:
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MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg

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RESULT 11
AAN81111
ID AAN81
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XX
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AAZ07505/c
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                                                                                                                                                                      US-09-857-841-4 (1-14) x AAZ07505 (1-1067)
                                                                                                                                                                                                                                                                                    Alignment Scores:
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                             AAN81111;
                                                                                                                                                                                                                                                                                                                Sequence 1067 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Preparing transgenic plants resistant to RNA virus infection - usi anti:sense gene constructs containing the viral coat protein gene,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-MAY-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ07505 standard; cDNA; 1067
                                                        AAN81111 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Fig 1; 15pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           from cucumber mosaic virus.
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69.00
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25-FEB-1987;
18-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                   This DNA is produced on screening of a plasmid library and is used to transform plant cells which subsequently produce the coat prote ein of cucumber mesaic virus (CMV) strain Y. This protein is not synthesised natural plant cells. The resistance to CMV infect ion, provided by the DNA, is shown in plant cells and redifferent iated plant bodies. See also AAN81110. (Updated on 25-MAR-2003 to correct PR field.) (Updated
Cucumber mosaic virus; tomato mosaic virus; minus virus resistance; disease resistance; transgenic
                                                       25-MAR-2003
20-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA coding for the coat protein of cucumber mosaic virus strain Y - used for producing plants resistant to cucumber mosaic virus infection.
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12-NOV-1990
                                 Cucumber mosaic virus RNA-3 chimera encoding RNase T1.
                                                                                                                AAQ76107 standard;
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(first entry)
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87JP-00043443.
88JP-00035809.
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           minus-strand RNA;
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plant; cross protection;
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Percent Similarity:
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                                                                                                                                                                                                                                                                                                                                        Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                       A chimeric cucumber mosaic virus RNA-3 (given in AAQ76107) has the coat protein gene replaced by a gene encoding a cell inhibitory protein, RNase T1, having the sequence given in AAR67754. The construct elicits a minussense RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus-resistance on a host plant, e.g. tobacco, tomato. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA constructs for imparting pathogen, esp. virus, resistance to plants encodes RNA that interacts with viral RNA polymerase to generate an eliciting agent.
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20-JUL-1995
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cell inhibitory
                      hypersensitive
                                   virus resistance;
                                             Cucumber mosaic virus; tomato mosaic virus; minus-strand RNA;
                                                                    Cucumber mosaic virus RNA-3 chimera encoding ToMV CP
                                                                                                                             AAQ76106;
                                                                                                                                                     AAQ76106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cucumber
            Lycopersicon
                                                                                                                                                                                                                                                                                                                                No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-DEC-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hypersensitive response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1995-036490/05.
DB; AAR67754.
                                                                                                                                                                                                   733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SANDOZ LTD.
SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P
                                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 31-32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mosaic
                                                                                                                                                                                                  ATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGA
                                                                                                                                                                                                               MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                esculentum;
ory protein;
                                                                                           (revised)
(first entry)
           ce; disease resistance; transgenic p
response; crop improvement; tomato;
sculentum; tobacco; Nicotiana tabacu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     93GB-00011593
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94WO-EP001817
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers 123. .437
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/product= "RNase T1"
                                                                                                                                                   DNA;
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69.00
100.00%
100.00%
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                                                                                                                                                     1860
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tobacco; Nicotiana tabacum; RNase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOMV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VERW
                                                                                                                                                     BP
                                                                                                                                                                                                                                                                                                                                                                 C; 389 G; 507
                                                                                                                                                                                                                                               (1-1696)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CMV; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GES
                                                                                                                                                                                                                                                                               Length:
Matches:
Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                 H,
            tabacum;
                                                                                                                                                                                                                                                                                                                                                                 0 U; 0 Other;
                                                                                                                                                                                                                                                                     1696
0
0
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                                 plant; cross protection,
                                                                                                                                                                                                   774
                                                                                                                                                                                                                        14
            coat
           protein;
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            CP;
                                                                                                                                                              RESULT 14
AAQ76108
                                                                                                                                                                                                                                                                              Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                 US-09-857-841-4 (1-14)
                                                                                                                                                                                                                                                                                                                                           Alignment
                                                                                                                                                                                                                                                                                                                                                                                       A chimeric cucumber mosaic virus RNA-3 (given in AAQ76106) codes for the coat protein (CP) of tomato mosaic virus (AAR67752) as well as its own CP (AAR67753). The construct elicits minus-sense RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus-resistance on a host plant, e.g. tobacco, tomato. (Updated on 25-MAR-2003 to correct PN field.)
        virus resistance; disease resistance; transgenic plant; cross protection; hypersensitive response; crop improvement; tomato; Lycopersicon esculentum; tobacco; Nicotiana tabacum; P30; elicitor; ToMV;
                                                                                          25-MAR-2003
20-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             eliciting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA constructs for imparting pathogen, encodes RNA that interacts with viral I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SANO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CDS
                                             Cucumber mosaic virus; tomato mosaic virus; minus-strand RNA;
                                                                   Cucumber mosaic virus RNA-3 chimera encoding
                                                                                                                                                   AAQ76108
                                                                                                                                                                                                                                                                                                                                                                 Sequence 1860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1995-036490/05.
DB; AAR67752, AAR67753
                                                                                                                                                                                                   897
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SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page
                                                                                                                                                                                                 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               agent.
                                                                                          (revised)
(first en
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English.
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Matches:
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                 552 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        esp. virus, resistance to plants
RNA polymerase to generate an
                                                                     TOMV
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                                                                                                   24-OCT-2003
27-AUG-2003
25-MAR-2003
07-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                                    A chimeric cucumber mosaic virus RNA-3 (given in AAQ76108) has the coat protein gene replaced by a gene encoding an elicitor, TOMV P30, having the sequence given in AAR67755. The construct elicits a minus-sense RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus-resistance on a host plant, e.g. tobacco, tomato. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA constructs for imparting pathogen, esp. virus, resistance to encodes RNA that interacts with viral RNA polymerase to generate eliciting agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SANO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CDS
                                                          CMV-C;
                                                                               Cucumber Mosaic
                                                                                                                                                                                                                                                                                                                                                                  Sequence 2173 BP; 561 A; 438 C; 532 G; 642 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1995-036490/05.
P-PSDB; AAR67755.
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                                       Cucumber mosaic
                                                                                                                                                       AAQ03641;
                                                                                                                                                                            AAQ03641 standard; DNA; 657
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 5; Page 34-35; 50pp; English.
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                   WO9002185-A.
                                                                                                                                                                                                                                                                                                                                    No.:
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                                                                                                                                                                                                                    1210 ATGGACAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGA 1251
                                                          curcurbitaceae; solanaceae; ds
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SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN VERW GES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PT;
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(first entry)
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                                       virus;
                                                                               Virus-C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= a
/product= "ToMV
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69.00
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                                       strain C
                                                                             (CMV-C) coat protein gene.
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                                    US-09-857-841-4 (1-14) x AAQ03641
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                                                                                                                                                 The sequence encodes the coat protein of CMV-C. (Updated on 25-MAR-2003 to correct PF field.) (Updated on 25-MAR-2003 to correct PA field.) (Updated on 27-AUG-2003 to correct OS field.) (Updated on 24-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                 Coat protein gene of cucumber mosaic transformed plants which are resistar
                                                                                                                                Sequence 657 BP;
                                                                                                                                                                                                 Disclosure; Page ?; 18pp; English.
                                                                                                                                                                                                                                                WPI; 1990-099409/13.
                                                                                                                                                                                                                                                                  Quemada H,
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Slightom JL,
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                                                               0.0379
64.00
92.86%
92.86%
92.75%
                                                                                                                                                                                                                    are resistant
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                                     (1-657)
                                                    Length:
Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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                                                                                                                                                                                                                   virus strain WL - cloned
nt to CMV viral infection
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Search completed: December Job time : 329 secs 08:24:35

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Result
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-MODEL-frame+ p2n.model -DEV=xlh
-Q-/cgn2_1/USPTO_spool/US08857841/runat_03122004_160551_18593/app_query.fasta_1.199
-DB=EST -QFMT=fastap -SUFFIX=p2n.rst -MINMATCH=0.1 -LOOFCL=0 -LOOFEXX=0
-UNITS-bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS-bluman40.cdi -LIST=45
-DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE=LOCAL
-OUTFWT=pto -NORM-ext -HEAPSIZE=500 -WINLEN=0 -MAXLEN=200000000
-USER-US09857841_0CGN 1 1 3437 @runat_0312204 160551_18593 -NCPU=6 -ICPU=3
-NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -DSSPECOK=100 -LONGLOG
-DEV TIMEOUT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPDXT=0.5 -FGAPOP=6
-PGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Perfect score:
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Listing first 45 summaries
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Maximum DB
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BI971551 GM830013B
AQ737846 HS 2157 A
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AL3	AQ8	BQ1	AJ5	AW1	AZO	AV1	BF5	BG40	900	CC4	BU2	CF2	CEO	CFR	CB2	AI9	800	AW7	ð	BM542:	BU13	BU32	BU2	BU25	BU2	BU3	ဌ	OE6	BU3	AG5	CF.	CD3	BX2	BX2	AL8
59	6379	64	2	2	494	61	15	0405	2	0	71	u	15692	8	4	۳	9	5242	702						2	õ	õ	ŭ	4	4	7	560	567	567	362
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A	AQ8	뽔	٤	A	ΑŽ	Ą	<b>H</b>	B B	ဥ	ဥ	먪	끉	В	끉	В	ΑI	გ	A	A	무	胺	BC	瞪	BG	명	BC	ВС	В	멷	ĕ	G G	CD3	æ	B	AL8
		BQ164	502	1127	049	161	519	404	692	401	1271	250	056	881	241	930	.098	1752	770	1542	1134	BU323014	1227	1256	1261	1366	1109	629	1340	540	577	356	BX256	BX256	w
661	3793					127	533	055	105	658	993	879	926	443	923						521	014	211	140	839	212	941	751			172		6742	w	4
MtB	nbeb	EST	AJ50:	M11	GSSI	-	EST	602	OGWDU6	HDA	6038	8	tig	tric	HI-	u160	NDL.	RC6	SH	AGE	603	603	0	2	603	8	8	ï.	603	Mus	MCS	AGE	BX25	BX2	AL8
BA32E	002	901	291	0634	Bruo	61127	1569	4199	9069	09 IMHD	ᇎ	Ħ	r-gas	2	ን	0e09	.107K1	-CT02	2260	BNCOURT	312103	40609	1339948	7459	150378	7890	1266	6-3a	191	musc	MCSA209G	AGENCOUR!	574	574	520
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# ALIGNMENTS

source	FEATURES	JOURNAL COMMENT	TITLE	REFERENCE	ACCESSION VERSION KEYWORDS SOURCE ORGANISM	RESULT 1 BM067231/c LOCUS DEFINITION
1127 /organism="Capsicum annuum" /mol_type="mRNA" /cultivar="Hang Keun" /db_xref="taxon:4072"	Korea Research Institute of Bioscience and Biotechnology P.O. Box 115, Yusong, Taejeon, 305-600, Republic of Korea Tel: 82-42-860-4340 Fax: 82-42-860-4309 Email: doil@mail.kribb.re.kr High quality sequence stop: 127. Location/Qualifiers	Annual L.) and bequeine Analysis in Relation to Hypersensitive Response Against Pathogen Unpublished (2001) Contact: Doil Choi Choi Choi Choi Choi Choi Choi Choi	Hur(CG. and Choi, D.  Generation of Expressed Sequence Tags from Hot Pepper (Capsicum Generation of Expressed Sequence Tags	Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids; lamids; Solanales; Solanaceae; Capsicum.  1 (bases to 127)  1 (bases Kim S - V Chung V - 4 Shin H - 1 Gob S - 4 Dai H - 5	EM067231 EM067231.1 GI:22787366 EST. Capsicum annuum Capsicum annuum	BM067231 127 bp mRNA linear EST 11-SEP-2002 KS08001E12 KS08 Capsicum annuum cDNA, mRNA sequence.

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RESULT 2
BE023342/c
LOCUS
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Best Local Similarity:
Query Match:
DB:
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                                                                                                                                                                                                                                                                                                                                              source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         #16 bp mRNA linear EST 24-JUL-2004 sm70f09.yl Gm-c1028 Glycine max cDNA clone GENOME SYSTEMS CLONE ID: Gm-c1028-9282 5' similar to TR:013437 Q13437 DELETED IN SPLIT HAND/SPLIT FOOT 1 ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              78
                                                                                                                                                                                                                                                                                                                                                               When it has been determined, an EST from the other end of this clone is listed in the 'Other ESTs on clone' field. This clone is available through: Biogenetic Services, 801 32nd Ave. Brookings, SD 57006 USA (phone: 800 423 4163; email: info@biogeneticservices.com) Insert Length: 1035 Std Error: 0.00.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Other ESTs: B1971551 corresponding to Gm-
Contact: Shoemaker R/Public Soybean EST I
Public Soybean EST Project
Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shoemaker, R., Keim, P., Vodkin, L., Erpelding, J., Coryell, V., Robana, A., Bolla, B., Marra, M., Hillier, L., Kucaba, T., Martin, J., Kecke, C., Wylie, T., Underwood, K., Steptoe, M., Theising, B., Allen, M., Bowers, Y., Person, B., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., Ritter, E., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R. and Wilson, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; spermatophyta; Magnoliophyta; Eudicotyledons; core eudicots; rosids; eurosids I; Fabales; Fabaceae; Papilionoldeae; Phaseoleae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EST.
Glycine max (soybean)
Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Public Soybean EST Project
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BE023342.1 GI:8285783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (bases 1 to 416)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Forest Park Parkway, Box 8501, St. Louis,
                      'Supernod' plants generously donated by Dr. Gary Stacey. The seedlings were innoculated with Bradyrhizobium japonicus, strain USDA110 priot to harvest. Stratagene's
                                                                                           /note="Vector: pBluescript II XR; Site 1: EcoRI; Site 2: XhoI; The mRNA was isolated from roots of Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="anther"
/dev_stage="10 weeks after germination"
/clone lib="K808"
/note="Vector: pBluescript SK(-)"
                                                                                                                                                                  db xref="taxon:3847"
/clone="GENOME SYSTEMS CLONE ID: Gm-c1028-9282"
/tissue_type="roots of 'Supernod' plants"
/tab host="DHIOB"
japonicus, stra
DNA synthesis
                                                                                                                                                clone_lib="Gm-c1028"
                                                                                                                                                                                                                                                                       /mol_type="mRNA"
/cultivar="Supernod"
                                                                                                                                                                                                                                                                                                                    organism="Glycine max"
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69.00
100.00%
100.00%
Kit (catalog number 200401) was used to
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Matches:
Conservative:
Mismatches:
Indels:
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US-09-857-841-4 (1-14) x BE023342
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                                                                                                                    Public Soybean EST Project
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. J
                                                                                                                                                                                                                                                Shoemaker, R., Keim, P., Vodkin, L., Erpelding, J., Coryell, V., Khanna, A., Bolla, B., Marra, M., Hillier, L., Kucaba, T., Martin, J., Khanna, A., Wylie, T., Underwood, K., Steptoe, M., Thebing, B., Allen, M., Bowers, Y., Person, B., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., Ritter, E., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., Scann, R., Waterston, R., and Wilson, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CA800884.1 GI:26057970
EST.
Glycine soja
Glycine soja
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CA800884 443 bp mRNA linear EST 01-JUL-2004 sat26b02.yl Gm-c1056 Glycine soja cDNA clone SOYBEAN CLONE ID: Gm-c1056-2379 5' similar to TR:Q9XIR8 Q9XIR8 F13011.6 PROTEIN. ;,
When it has been determined, an EST from the other end of this clone is listed in the 'Other ESTs on clone' field. This clone is available through: Biogenetic Services, 801 32nd Ave. Brookings, SD
                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
                                                                Email: est@watson.wustl.edu
                                                                                     Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                        Contact: Shoemaker R/Public Soybean EST Project
                                                                                                                                                                                                            Unpublished (1999)
                                                                                                                                                                                                                                 Public Soybean EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATGGACAAAGCCGAATCTACCGGCTGTGGAAGAACGAAAAGA
                                                                                                                                                                                                                                                                                                                                                                               (bases 1 to 443)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   synthesize the cDNA. First-strand synthesis was with 5-methyl dCTP, hence the ligated cDNA was hemimethylated. A modification of Stratageness hemimethylated.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92.7
49.00
78.57%
64.29%
71.01%
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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                                                                                                                            St. Louis, MO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2
3
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                                                                                                                                63108, USA
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Score: Alignment ORIGIN

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TITLE JOURNAL COMMENT

REFERENCE AUTHORS

ACCESSION VERSION

SOURCE

ORGANISM

KEYWORDS

LOCUS DEFINITION

CA800884/c RESULT 3

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TITLE
JOURNAL
COMMENT
                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE
AUTHORS
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Query Match:
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BI468772/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM
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456 bp mRNA linear EST 08-JUL-2004 sai03c10.y1 Gm-c1050 Glycine max cDNA clone GENOME SYSTEMS CLONE ID: Gm-c1050-4363 5' similar to TR:Q9XIR8 Q9XIR8 F13011.6 PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Glycine max
                                                                                                                                                                                                                                              Shoemaker, R., Keim, P., Vodkin, L., Erpelding, J., Coryell, V., Khanna, A., Bolla, B., Marra, M., Hiller, L., Kucaba, T., Martin, J., Beck, C., Wylie, T., Underwood, K., Steptoe, M., Theising, B., Allen, M., Bowers, Y., Person, B., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., Ritter, E., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R. and Wilson, R.
Public Soybean EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Spermatophyta; Magnoliophyta; eurosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Glycine max (soybean)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BI468772.1 GI:15284890
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                                                                                                        Washington University School of Medicine
                                                                                                                                                               Contact: Shoemaker R/Public Soybean EST Project
Public Soybean EST Project
                                                                                                                                                                                                                         Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATGGACAAAGCCGAATCTACCGGCTGTGGAAGAACGAAAAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                             (bases 1 to 456)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone lib="Gm-cl056"
/clone lib="Gm-cl056"
/clone lib="Gm-cl056"
/clone lib="Gm-cl056"
/clone lib="Wector: pBluescript II SK+; Site 1: EccRI; Site 2:
/note="Wector: pBluescript II SK+; Site 1: EccRI; Site 2:
/note="Wector: pBluescript II SK+; Site 1: EccRI is olated
from 4 day old seedling of p1468916. The seedlings were
germinated in a growth chamber using germination paper.
Complementary DNA was synthesized from mRNA using a primer
consisting of a poly(dT) sequence with a XhoI restriction
site. EccRI adapters were ligated to the blunt-ended cDNA
fragments followed by XhoI digestion. The cDNA fragments
were directionally cloned into the EccRI-XhoI restriction
site of the pBluescript vector. The ligated cDNA fragments
were transformed into DH10B host cells (GibcoBRI). This
'ihrarv was constructed in the laboratory of Dr. Randy
                                                    314 286 1800
314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           quality sequence stop: 421.
  has been determined,
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/tissue_type="Whole seedling, 4 day old"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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49.00
78.57%
64.29%
71.01%
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Matches:
Conservative:
Mismatches:
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RESULT 5
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                                J (bases 1 to 544)
Vodkin, L., Keim, P., Shoemaker, R., Retzel, E., Khanna, A., Erpelding, J., Raph, C., Shoop, E., Pardinas, J., Liu, L. and A Functional Genomics Program for Soybean (NSF 9872565)
Unpublished (1999)
Other_ESTs: BE023342 corresponding to Gm-c1028-9282 (5')
Contact: Vodkin, L.O., PI, A Functional Genomics Program Soybean (NSF 9872565)
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                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
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GM830013B21D12 Gm-r1083 Glycine max cDNA clone Gm-r1083-4800 3',
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Lewin, H. A., Director, Keck Center for Comparative and Functional
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/tissue_type="leaf tissue at various developmental stages
of greenhouse grown plants"
/dev_stage="3 week old"
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'cultivar="Clark"
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RESULT 6
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HS_2167_A1_B08_T7C CIT Approved Human Genomic Sperm Library D Homo
sapiens genomic clone Plate=2167 Col=15 Row=C, genomic survey
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                                          Homo
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                                                                                                   AQ737846.1
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Reracking was performed by Incyte Genomics, St. Louis, http://www.incyte.com, and 3' sequencing by the Keck Center for Comparative and Functional Genomics, University of Illinois, http://www.lie.uiuc.edu/biotech/keck.html.

Note: The corresponding 5' EST from each clone in the detailed information on the source library for each clone can also be obtained by referring to the Incyte Genomics clone clone ID of the original CDNA library that is also listed under 'OTHER EST',"
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Medical Center Drive, Rockville, MD 20850, USA

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              clade; Panicoideae; Andropogoneae; Zea.

(bases 1 to 956)

Whitelaw.C.A., Quackenbush, J., Van Aken, S., Utterback, T.,
Resnick, A., Fraser, C.M., Budiman, M.A., Bedell, J.A., Rohlfing, T.,
Citek, R.W., Nunberg, A., Robbins, D. and Lakey, N.

Consortium for Maize Genomics
Unpublished (2002)
Other GSSs: ZUAG628TV
Other GSSs: ZUAG628TV
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ZUAG628HH ZM 3.0_4.0 KB Zea
genomic survey sequence.
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1 (bases 1 to 617)

Mahairas,G.G., Wallace,J.C., Smith,K., Swartzell,S., Holzman,T.,
Mahairas,G.S., Wallace,J.C., Young,J., Zhao,S., Adams,M.D. and
Keller,A., Shaker,R., Furlong,J., Young,J., Zhao,S., Adams,M.D.
                                                                                                                                                                             Zea mays
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD
                                                                                                                                                                                                                                                                 GSS.
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High quality sequence stop:
Location/Qualifiers
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Contact:
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Clones may be purchased from Research Genetics (info@resgen.com).
BAC end Web Server: http://www.htsc.washington.edu
Plate: 2167 row: C column: 15
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401 Queen Anne Avenue North, Seattle,
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/note="Organ: sperm; Vector: pBeloBAC11; BAC Clones i
E-Coli DH10B"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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ORIGIN
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                                                                                                                                                                                                                                                                                   Contact: Jiang Lu, FAMU, Wayne Hunter, US
Viticulture Centre, FAMU, Tallahassee, FI
Florida A&M University
Tallahassee, FI 32317, USA
Tel: (850) 412-7393
Fax: (850) 561-2617, (772) 462-5898
Email: jiang.lu@famu.edu,
Seg primer: T3 Primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  213 CTGGACAAGACGAAGCAGACCGCCGTGGGAAGAAACCGGCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                  Hunter, W.B., Dang, P.M., Chaparro, J.X., Lu, J. and Leong, S. Genes expressed in Vitis shuttleworthii L Unpublished (2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; Vitaceae; Vitis.
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CF568862
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Class: sheared ends.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vitis shuttleworthii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CF568862.1 GI:34994945
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       /sex="Mixed population"
/tissue_type="Entire tendril, leaves, bud, flowers"
/dev_stage="At blooming"
/lab_host="XLI-Blue"
/clone_lib="Vitis shuttleworthii L., grape"
/note="Vector: pBluescript II SK+; Site 1: EcoRI; Site 2:
XhoI; A high quality EST with at least 100 contiguous bases at Trace Tuner score of 20 or better. Construction by PM Dang, USDA, ARS, U.S. Horticultural Research Lab, Ft.Pierce, FL, USA."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              genomic DNA library"
                                                                                                                                                                                              /organism="Vitis shuttleworthii"
/mol_type="mRNA"
/db_xref="taxon:246827"
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/note="Vector: pBCSK-; Site_1: HincII; 3-4 kb 'unfiltered'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Zea mays"
/mol_type="genomic DNA"
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/clone="ZMMBPa0057F07"
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48.00
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REFERENCE
AUTHORS
TITLE
RESULT 10
AL836204
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AL584972/c
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                                               267 ATACAGAAATCTGAAAGTACAGCTCCTGGAAACAACAGGCGG 226
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AL584972
AL584972.1 GI:13163705
EST.
Gallus gallus (chicken)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                73 ACAGAGACAACATCTGCAGGGAGAAATCGGAGA 105
                                                                       1 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg 14
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Roslin, Midlothian, EH25 9PS, UK
Tel: +44 (0)131 527 4200
Pax: +44 (0)131 440 0434
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Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
Phasianinae; Gallus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: frazer.murray@bbsrc.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Frazer Murray
Dept. Genomics and Bioinformatics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Stratagene Chick Embryo Lambda cDNA Library
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                                                                                                                                                                                                                                                                                                                                                                             /dev_stage="5 days old"
/lab_host="SOLR cells (kanamycin resistant)"
/clone lib="Stratagene Chick Embryo Lambda cDNA Library
937405)"
                                                                                                                                                                                                                                                                                                            /note="Vector: pBLUESCRIPT SK; Site_1: BcoR1; Site_2: Xho1; Cloned unidirectionally. Primer: Oligo dT. Uni-ZAP XR vector. Average insert size: 1.5kb; 5' adaptor sequence: 5' GAATTCGGCACGAG 3'; 3' adaptor sequence: 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Gallus gallus"
/mol_type="mRNA"
/db_xref="taxon:9031"
/clone="ROS016G12"
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Chick Embryo Lambda cDNA Library (* 937405)
clone ROS016G12, mRNA sequence.
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BX256743
Gallus gallus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
Phasianinae; Gallus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MRC Human Genome Mapping Project Resource Hinxton, Cambridge, CB10 15B, UK Email: biohelp@hgmp.mrc.ac.uk Email: biohelp@hgmp.mrc.ac.uk Vector: pBluescript II KS
                                                                    Gallus gallus (chicken)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 480)
Clark,M.S., Edwards,Y.J., Peterson,D., Clifton,S.W., Thompson,A.J. Sasaki,M., Suzuki,Y., Kikuchi,K., Watabe,S., Kawakami,K., Sugano,S., Elgar,G. and Johnson,S.L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AL836204 EFRd Takifugu rubripes cDNA clone EFRd002apsA11, mRNA
                                                                                                        BX256743.2
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Cambridge, CB10 1SB. UK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genome Res. 13 (12), 2747-2753 (2003)
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BX256743 AGENAE Gallus gallus multi-tissues normalized library
(gcag) Gallus gallus cDNA clone gcag0001c.c.05 5prim, mRNA
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PRIMER: KS
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                                                                                                                                                                                                                                                                                    AspLysSerGluSerThrSerAlaGlyArgAsnArgArg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="gut"
/clone_lib="EFRd"
/note="Vector: pBlv
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/mol_type="mRNA"
/db_xref="taxon:31033"
/clone="EFRd002apsA11"
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BX256742
   Construction
                                                                                                                         Gallus gallus
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                                                                                                                                                                             BX256742.2
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US-09-857-841-4 (1-14) x BX256743 (1-486)
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                                                                                                                                                                                                                                                                                                                      BX256742 AGENAE Gallus gallus multi-tissues normalized lib (gcag) Gallus gallus cDNA clone gcag0001c.c.05 3prim, mRNA
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Herault,F., Le Meuth-Metzinger,V., Desert,C., Retout,E., Piumi,F.,
Klopp,C. and Douaire,M.
Construction and primary characterization of chicken normalized
multi-tissue cDNA libraries
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
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Sequence cleaned of vector, adaptator and repetitions. Co
at sigenasupport@jouy.inra.fr to obtain the chromatogram
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65, rue de Saint-Brieuc, RENNES cedex, 35042,
Tel: +33 (0) 2.23.48.54.63
Fax: +33 (0) 2.23.48.54.70
                     1 (bases 1 to 490)
Herault, F., Le Meuth-Metzinger, V., Desert, C.,
Klopp, C. and Douaire, M.
                                                                                            Archosauria; Aves; Neognathae; Phasianinae; Gallus.
                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
                                                                                                                                                                                                Gallus gallus (chicken)
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/tissue_type="multi-tissues"
/dev_stage="from embryos to adults"
/lab_host="DH108"
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|mol_type="mRNA"
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and primary characterization of chicken normalized
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                                               Retout, E.,
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                                                  Piumi, F
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REFERENCE
AUTHORS
TITLE
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CD356033
LOCUS
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KEYWORDS
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Tel: +33 (0) 2.23.48.54.63

Fax: +33 (0) 2.23.48.54.70

Emmil: Madeleine.Douaire@roazhon.inra.fr

Sequence cleaned of vector, adaptator and repetitions. Contact us at sigenasupport@jouy.inra.fr to obtain the chromatogram of this
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AGENCOURT 14292133 NIH MGC 176 Mus
IMAGE:30379119 5', mRNA sequence.
CD356033
                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleosto
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
1 (Dases 1 to 645)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                      Contact: Daniela S. Gerhard, Ph.D. Office of Cancer Genomics National Cancer Institute / NIH Bldg. 31 Rm10A07 Bethesda, MD 20892
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On Feb 27, 2003 this sequence version replaced gi:28579340.
Contact: Douaire M
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CD356033.1 GI:31127444
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                                                                                                                                                                                                                               Unpublished (1999)
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cgapbs-r@mail.nih.gov
Procurement: Dr. Michael Brownstein
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library (gcag)"
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/clone="gcag0001c.c.05"
/tissue_type="multi-tissues"
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/mol_type="mRNA"
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/lab_host="DH10B"
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JOURNAL COMMENT
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Bonnett, G.D. and Manners, J.M.
Identification of differentially expressed transcripts from maturing stem of sugarcane by in silico analysis of stem expressed sequence tags and gene expression profiling Unpublished (2003)
Commonwealth Scientific and Industrial Research Organisation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Saccharum.
                                                                                                                                                                                             Division of Plant Industry Long Pocket Laboratories,
                                                                                                                                                                                                                                                                                                                                                                                                                     Casu, R.E., Dimmock, C.M., Chapman, S.C., Grof, C.P.L, McIntyre, C.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 653)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Saccharum sp.
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Location/Qualifiers
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//clone="Iba" MAGE:30379119"
//lab host="DH10B (71-phage-resistant)"
//clone_lib="NIH MGC 176"
//clone_lib="NIH MGC 176"
//clone_lib="NIH MGC 176"
//clone_lib="Organ: kidney; Vector: pDNR-LIB; Site_1: SfiI
//cloned.5; cDNA made
by oligo-dT priming and directionally cloned.5; and 3;
adaptors were used in cloning as follows:
5; ARGCAGTGGTATCAACGCAGAGTGGCCATTACGGCCGGG-3; and
5; ARGCAGTGGTATCAACGCAGAGTGGCCATTACGGCCGGG-3; and
5; ATTCTAGAGGCCGAGGCGGCGACATG-dT(30)NN-3; Full-length
enriched library was constructed using the Clontech
Creator SMART kit and size-selected to contain the 0.5 kb
size fraction. Library created in the laboratory of M.
Brownstein (NIMH, NIH). Note: this is a NIH_MGC Library."
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/db_xref="taxon:10090"
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AUTHORS
TITLE
JOURNAL
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AUTHORS
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SOURCE
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Best Local Similarity:
Query Match:
DB:
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JOURNAL
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2 Thases 1 to 677)

RS Hattori,M., Toyoda,A., Noguchi,H., Kojima,T. and Sakaki,Y.

Direct Submission

AL Submitted (17-NOV-2003) Masahira Hattori, The Institute of Physical and Chemical Research (RIKEN), Genomic Sciences Center (GSC);

1-7-22 Suehiro-chou,Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan (E-mail:hattori@gsc.riken.jp, URL:http://hgp.gsc.riken.go.jp/,

Tel:81-45-503-9111, Fax:81-45-903-9170)

Clones are derived from the mouse BAC library MSMg01. For BAC library availability, please contact Kuniya Abe (abe@rtc.riken.jp).

The Institute of Physical and Chemical Research (RIKEN) 3-1-1 Koyadai, Tsukuba, 305-0074 Japan phone: 81-298-36-9189, fax: 81-298-36-9199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vector
R.Site 1
R.Site 2
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AG540710
                                                            Sequencing : TJ
LIBRARY
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                  Hattori,M., Toyoda,A., Noguchi,H., Kojima,T. and Sakaki,Y. BAC end Sequences of Library MSMg01
Unpublished
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GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone_lib="Maturing Sugarcane Stem Lambda ZIPLOX Library
(MCS)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /dev_stage="12 months post-planting"
/lab_host="Y1090(ZL)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:15819"
/clone="MCSA189F12"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cultivar="Q117"
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46.00
85.71%
64.29%
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Matches:
Conservative:
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clone:MSMg01-453M10.TJ,
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, genomic survey
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Percent Similarity:
Best Local Similarity:
Query Match:
Search completed: December Job time: 2049 secs
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                                                                                                                    US-09-857-841-4 (1-14) x AG540710
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                                                          284 ATGAACAAAAGGGAGCAAGTAGAGGCAGGCAGGAACAGAAGA 243
                                                                            1 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
                                                                                                                                                                                                                                                                                                                   /organism="Mus musculus molossinus"
|mol type="genomic DNA"
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|db_xref="reaxon:57486"
|clone="MSMg01-453M10.TJ"
|sex="male"
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                                                                                                                                                                                                                                                                                     /tissue_type="mixture of kidney and spleen"
/clone_lib="MSMg01 Mouse Male BAC Library"
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Result
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-DB=ISSUED_Ratents_NA -QPMT=fastap_SUPFIX=ppn.rni -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=51ts -GTART=1 -END=-1 -MATRIX=b10sum62 -TRANS=human40.cdi
-LIST=45 -DCCALLGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pco -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER=US09857841 @CGN 1 1 69 @TUNATE 03122004 160552_18608 -NCPU=6 -ICPU=3
-NO MMAP -LARGEQUERY -NEG_SCORES=0 -MAIT -DSPBLOCK=100 -LONGLOG
-DSV_TIMEOUT=120 -MARM_TIMEOUT=30 -THREADS=1 -XOAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 2000000000
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Ygapop 10.0 , y
Fgapop 6.0 , F
Delop 6.0 , F
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/cgn2 6/ptodata/1/ina/5B_COMB.seq:*
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148.523 Million cell updates/sec
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Fgapext
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US-08-488-031-14
US-08-488-059-14
US-08-488-059-14
US-08-090-192-14
US-08-090-192-14
US-08-482-663-14
US-08-482-658-14
US-08-475-610-14
US-08-475-610-14
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Appl	Appl	App1	Appl	ildö	1dd	Appli	1gg	pli	lgg^	ppli	φp1	opli	App1	ppli	1qqA	ppli	App1	ppli	1qqA	ppli	1gg/	ppli	Appl	App1	App1	App1	App1	App1	App1	App1	Appl	App1

### ALIGNMENTS

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equence 14, Application US/08090193
atent No. 5641673
APPLICANT: Haseloff, James
APPLICANT: Brand, Andrea
APPLICANT: Brand, Andrea
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
ADDRESSE: Sterne, Kessler, Goldstein & Fox
STREET: 1100 New York Avenue, Suite 600
CITY: Washington
STATE: D.C.
COUNTRY: U.S.A.
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATIOS SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
APPLICATION DATA:
APPLICATION NUMBER: US/08/090,193
FILING DATE: 23-DEC-1993
CLASSIFICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,330
FILING DATE: 1-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
REGISTRATION NUMBER: 33,851
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APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: BUGBISKY, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080002/MAC/LBB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2540
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 130 base pairs
TYPE: nucleic acid
Grean Application of the control of the co
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,031
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,193
FILING DATE: 23-DEC-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Haselo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS: ADDRESSE: Sterne. Konnik
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 130 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 20005
COMPUTER READABLE FORM:
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
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Matches:
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Percent Similarity:
Best Local Similarity:
Query Match:
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DB:
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                                                                                    Alignment Scores:
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Best Local Similarity:
                                                                    Pred. No.:
                                                                                                                     US-08-486-569-14
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Patent No. 5
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FILING DATE: 07-UN-1995
PRIOR APPLICATION DATA:
APPLICATION UMBER: 08/090,193
FILING DATE: 23-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-DAN-1992
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
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APPLICANT: Haseloff, James
                                                                                                                                                                                                                        TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAW-1991
ATTORNEY/AGENT INFORMATION:
NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080006/MAC/LBB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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APPLICANT: Perrimon, No. 5863774bert
APPLICANT: Gerrimon, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 130 base pairs
                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
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TOPOLOGY: lir
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CITY: Washington
                                                                                                                                                                       TYPE:
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5863774
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Perrimon, No. 5863774bert
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Conservative:
Mismatches:
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Matches:
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RESULT 4
US-08-488-027-14
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RESULT 5
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Best Local Similarity:
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Patent No. 5866384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PCT/US92/00277
APPLICATION DATA: 16-JAN-1992
PRIOR APPLICATION DATA: US 07/642,330
APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (202) 371-2500
TELEPAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
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FILING DATE: 07-UN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,193
FILING DATE: 23-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/00277
APPLICATION NUMBER: PCT/US92/00277
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MEDIUM TYPE: Floppy disk
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TELECOMMUNICATION INFORMATION:
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APPLICANT: Perrimon, No. 5866384bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
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                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: both
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                                       ATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGA 94
                                                               MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg 14
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Matches:
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Best Local Similarity:
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                                                                                                                                                                      Sequence 14, Application US/08482663
Patent No. 5882907
GENERAL INFORMATION:
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APPLICANT: Haseloff, James
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 51
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 07/642
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAMB: Cimbala, Michele A.
REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0609
                                                  APPLICANT: Haseloff, James
APPLICANT: Brand, Andrea
APPLICANT: Perrimon, No. 5882907bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                  CORRESPONDENCE ADDRESS: ADDRESSEE: Sterne, F
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,333
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PRIOR APPLICATION NUMBER:
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MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: both
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3: Sterne, Kessler, Goldstein & Fox
1100 New York Avenue, N.W., Suite 600
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Matches:
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COUNTRY:

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US-08-482-658-14
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Patent No. 6010904
GENERAL INFORMATION:
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FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA: 08/090,193
APPLICATION NUMBER: 08/090,193
FILING DATE: 23-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/002
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 07/64
PILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox STREET: 1100 New York Avenue, N.W., Suite 600
                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                             APPLICANT: Perrimon, No. 6010904bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                                         APPLICANT: Haseloff, James APPLICANT: Brand, Andrea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
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STRANDEDNESS: both
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                        COUNTRY: U.S.A.
                                                                                                                                               STATE:
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                                                                                                        ZIP:
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Matches:
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Indels:
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Percent Similarity:
Best Local Similarity:
Query Match:
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APPLICANT: Haseloff, James
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 51
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                                                  APPLICATION NUMBER: US/08/470,349
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,192
FILING DATE: 11-NOV-1993
APPLICATION NUMBER: PCT/ US 92/00278
                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-DAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
FILING DATE: 16-JAN-1992 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 07
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MEDIUM TYPE: Floppy
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APPLICATION NUMBER:
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TYPE: nucleic acid
STRANDEDNESS: both
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ZIP: 20005
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(202) 371-257
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 US 07/642,333
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Matches:
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RESULT 9
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Best Local Similarity:
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REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0609.3030001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
TELEPAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 130 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-857-841-4 (1-14) x US-08-470-349-14 (1-130)
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GENERAL INFORMATION:
                                APPLICATION NUMBER: 08/090,193

PILING DATE: 23-DEC-1993

PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/00277

PILING DATE: 16-JAN-1992

PRIOR APPLICATION NUMBER: US 07/642,330

PILING DATE: 17-JAN-1991

APPLICATION NUMBER: US 07/642,330

PILING DATE: 17-JAN-1991

ATTORNEY/AGENT INFORMATION:
NAME: BUGAISKY, LAWTENCE B.
REGISTRATION NUMBER: 35,086

REFERENCE/DOCKET NUMBER: 0609.3080004/MAC/LBB

TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/475,610
FILING DATE: 07-UN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,193
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ADDRESSEE: Sterne, Kessler, Goldstein & Fox
STREET: 1100 New York Avenue, N.W., Suite 600
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MEDIUM TYPE: Floppy disk
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APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
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ATTORNEY/AGENT INFORMATION:
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STREET: 1100 New 1
CITY: Washington
STATE: D.C.
             TELEPHONE: (202)
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Query Match:
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PCT-US92-00277-14
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jorge A
REGISTRATION NUMBER: 29,021
REFERENCE/DOCKET NUMBER: 0609.3496604
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)833-7533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 130 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (202)833-8716
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 130 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 19920116
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
ADDITATION NUMBER: NC 07/642 220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
CONTROLL WITHOUT DATA:
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CITY: Washington
STATE: DC
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APPLICANT: Perrimon, Norbert
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                    TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
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Goodman, Howard
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                                                          Length:
Matches:
Conservative:
Mismatches:
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US-09-857-841-4 (1-14) x PCT-US92-00277-14 (1-130)

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PCT-US92-00278-14
                                                                                                                                      US-08-090-193-11
                                                                                                                                                        RESULT 12
                                                                                                                                                                                                                                                                                                          Query Match:
DB:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                       Alignment Scores:
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                                                                           Sequence 11, Applicat Patent No. 5641673
GENERAL INFORMATION:
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CURRENT APPLICATION DATA:
APPLICATION UNMBER: PCT/US92/00278
FILING DATE: 19920116
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,333
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jorge A
REGISTRATION NUMBER: 29,021
REGISTRATION NUMBER: 29,021
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TELEFAX: (202)833-8716
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)833-7533
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APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: Nucleic Acid
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: NUCLEIC ACID
STRANDEDNESS: Single
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STREET: 1225 Connecticut Avenue, N.W., Suite 300
CTTY: Washington
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                                                                                                                  Application US/08090193
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Haseloff, James
Brand, Andrea
Perrimon, No. 5641673bert
Goodman, Howard M.
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Matches:
Conservative:
Mismatches:
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Percent Similarity:
Best Local Similarity:
Query Match:
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FILING DATE: 23-DEC-1993
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JAN-1992
PRIOR APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
REGISTRATION NUMBER: 33.851
REFERENCE/DOCKET NUMBER: 0609.3080001
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                             Sequence 11,
Patent No. 5
                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Haseloff, James
APPLICANT: Brand, Andrea
APPLICANT: Perrimon, No. 58
APPLICANT: Goodman, Howard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/090,193
COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                        APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                    STREET: 1100 New CITY: Washington STATE: D.C.
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                                      COUNTRY: U
ZIP: 20005
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                                                                                                                             ADDRESSEE:
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5849548
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                                                                                                                                                                                                                     Brand, Andrea
Perrimon, No. 5849548bert
                                                       U.S.A.
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Matches:
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Mismatches:
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SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:

US/08/488,031

COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS

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US-09-857-841-4 (1-14) x US-08-488-031-11 (1-131)
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PAtentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/486,569
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,193
FILING DATE: 23-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 11, Application US/08486569 Patent No. 5863774
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: 08/090,193

FILING DATE: 23-DEC-1993

PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/US92/00277

FILING DATE: 16-JAN-1992

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Perrimon, No. 5863774bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid STRANDEDNESS: both
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     54 ATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGA
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Matches:
Conservative:
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Indels:
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-486-569-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 11, Approx No. 5866384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Haselof
APPLICANT: Brand,
APPLICANT: Perrimo
                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,027
FILING DATE: 07-UN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,193
FILING DATE: 23-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox
STREET: 1100 New York Avenue, N.W., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080006/MAC/LBB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                               PILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US (
                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 131 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
PCT/US92/00277
ATTORNEY/AGENT INFORMATION:
NAME: Bugaisky, Lawrence
                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/
FILING DATE: 16-JAN-1992
                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       54 ATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/08488027
                                                                                                                                                                                                                                                                                                                                                                   U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Perrimon, No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Brand, Andrea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (202)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Haseloff, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                      JMBER: US 07/642,330
17-JAN-1991
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Matches:
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Indels:
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Search completed: December 5, 2004, 09:48:23 Job time : 68 secs

Result

Query

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Command line parameters:

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-Q=/cgn2 1/USPTO_spool/US09857841/runat_03122004_160552_18628/app_query.fasta_1.199
-DB=Published Applications_NA -QFMT=fastap -SUFFTX=p2n.rnpb -MINMATCH=0.1
-LOOPCL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62
-TRANS=human40.cdd -LIST=45 -DOCALIGN=200 -THR SCORE=pct -THR MAX=100
-THR MIN=0 -ALIGN=15 -MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
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-NCPU=6 -ICPU=3 -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -DSPBLOCK=100
-LONGIJG -DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5
-FGAPOP=6 -FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Perfect score:
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Ygapop 10.0 , y
Fgapop 6.0 , I
Delop 6.0 , I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Published_Applications_NA:*

1: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*

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112:
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/cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*
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Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5, 2004, 08:17:52; Search time 350 Seconds (without alignments) 219.793 Million cell updates/sec
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Ygapext
Fgapext
Delext
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIE

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544	440 440 420	332 34 35 36	309 309 309	118 118 118 120 121 121 121 121 121 121 121 121 121	10 11 11 11 11 11 11 11 11 11 11 11	7654000
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	99900	22222	60.99 60.99	00000000000	2220000000000	100.0 100.0 100.0 100.0 92.8 69.6 68.1
523 563 573	3 2 2 5 2 3 4 6 8	9924007	1068 1058 1058	2377 2979 3105 3175 125439 125439 1414 414	000000000000	Tength 771 772 772 792 960 3048
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-10-425-115-79864 -10-424-599-31741 -09-918-995-3374	-10-741-601-57 -10-437-963-60 -10-437-963-89 09-294-093B-13 -10-425-115-13	-10-425-115-20 -10-305-720-12 -10-305-720-3 -10-001-843-35 -10-017-863-35 -10-437-963-93 -09-800-274-1	US-10-027-632-717 US-10-027-632-717 US-10-437-963-34410 US-10-424-599-113581 US-10-437-963-93102	-10-425-115-17 -10-425-115-17 -10-425-115-86 -10-425-115-17 -10-450-826-84 -09-987-899-23 -10-425-115-23 -10-437-963-15	-10-437-963-3120 -10-437-963-49610 -10-437-963-92187 -10-437-963-64563 -10-437-963-64563 -10-282-122A-2845 -10-282-122A-2845 -10-437-963-86635 -10-437-963-64807	US-10-011-033-5 US-10-011-033-1 US-10-011-033-1 US-10-011-033-3 US-10-011-033-9 US-10-011-033-9 US-10-425-115-176381 US-10-437-963-2533
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#### ALIGNMENTS

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US-10-011-033-5

US-10-011-033-5

Publication No. US20020124286A1

Publication No. US20020124286A1

GENERAL INFORMATION:

APPLICANT: Boeshore, Maury L

McMaster, J. Russell

Tricoli, David M

Reynolds, John F

Carney, Kim J

TITLE OF INVENTION: Plants Resistant to C Strains of Cucumber Mosaic Virus

NUMBER OF SEQUENCES: 15

CORRESPONDENCE ADDRESS:

ADDRESSE: APPLICATE: 2 Prudential Plaza, 180 N. Stetson, Suite 4700

STATE: 11inois

COUNTRY: US.A.

ZIP: 60089

COMPUTER READABLE FORM:
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
                                                                                                                                                                                                                                                                            RESULT 2
US-10-011-033-1
; Sequence 1, Application US/10011033
; Publication No. US20020124286A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                        ફ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-857-841-4 (1-14) x US-10-011-033-5 (1-771)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 771 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    . NO. :
                                                                                                                                   APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5400
TELEFAX: 312-616-5460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
CORRESPONDENCE ADDRESS:
ADDRESSEB: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
CITY: Chicago
STATE: Illinois
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /standard_name= "COAT PROTEIN" SEQUENCE DESCRIPTION: SEQ ID NO: 5:
                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                           1 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: CDS
LOCATION: 3..660
COTHER INFORMATION: /codon_start= 3
/function= "ENCAPSIDATES VIRUS RNA"
/product= "COAT PROTEIN"
/gene= "CP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN: V-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.00072
69.00
100.00%
100.00%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length:
Matches:
Conservative:
Mismatches:
Indels:
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14
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-011-033-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-UUN-1997
ATTORNEY_AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
TELEFONE: 312-616-5400
TELEFAX: 312-616-5400
INFORMATION FOR SEQ ID NO: 1:
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                             Sequence 14, Application US/10011033 Publication No. US20020124286A1 GENERAL INFORMATION:
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::
ZIP: 60089
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANALSM: Cucumber Mosaic Virus
INDIVIDUAL ISOLATE: V-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN FALEASE #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                            NUMBER OF SEQUENCES: 15 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                            TITLE OF INVENTION: Plants Resistant to C Strains Cucumber Mosaic Virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
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                                                                                            CITY: Chicago
STATE: Illinois
COUNTRY: U.S.A.
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TYPE: nucleic acid
STRANDEDNESS: single
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FILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                       ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: CDS
                                                                                                                                                                                                                                                                         McMaster, J. A. Tricoli, David M
                                                                                                                                                                                                                                                                     Reynolds, John Carney, Kim J
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Matches:
Conservative:
Mismatches:
Indels:
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RESULT 4
US-10-011-033-3
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Best Local Similarity:
Query Match:
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FILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMPUNICATION INFORMATION:
TELEPHONE: 312-616-5400
TELEPAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 14:
                                                    COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-No. US20020124286A1-2001
                                                                                                                                                                   NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE ROCKEY, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
CITY: Chicago
STATE: 11inois
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: cDNA HYPOTHETICAL: NO
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PILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 772 base pairs
TYPE: nucleic acid
STRANDENNESS: single
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US-10-011-033-9
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 312-616-5400
TELEFAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 3:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION NDATA:
APPLICATION UNMBER: US/10/011.033
FILING DATE: 13-NO. US20020124286A1-2001
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
                                                                                                                                                                                                                                                                                                                                                   Reynolds, John r
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
PILING DATE: 26-JUN-1997
ATTORNEY/ACENT INFORMATION:
NAME: Lisa V Mueller
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORIGINAL SOURCE:
ORGANISM: CUCUMBER MOSAIC VIRUS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 792 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 15 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE DESCRIPTION: SEQ ID NO:
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STATE: Illinois
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                                                                                                                                                                                                                            ZIP: 60089
                                                                                                                                                                                                                                               COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                  ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION:
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Matches:
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Score: Pred. No.:

ATTORNEY/AGENT INFORMATION: TELECOMMUNICATION INFORMATION:

NAME: Lisa V.

Mueller

TELEPHONE: 312-616-5400

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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yohua
APPLICANT: Cao, Yongwei
ITITE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
ITITE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 176381
LENGTH: 3048
TYPE: DNA
ORGANISM: Zea mays
FEATURE:
ORGANISM: Zea mays
FEATURE:
OTHER INFORMATION: Clone ID: MRT4577_92452C.1
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                                                               US-09-857-841-4 (1-14) x US-10-425-115-176381 (1-3048)
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US-10-425-115-176381
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Publication No. US20040214272A1
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                 1 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
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CTGGACAAGACGAAGCAGACCGCCGTGGGAAGAAACCGGCGG 1075
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TYPE: nucleic acid
STRANDEDNESS: single
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LOCATION: 1..
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Boukharov, Andrey A.
APPLICANT: Boukharov, Andrey A.
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 253
LENGTH: 609
TYPE: DNA
                                                                                                                                                                                            ; OTHER INFORMATION: Clone ID: PAT_MRT4530_102831C.1 US-10-437-963-3120
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; OTHER INFORMATION: Clone ID: PAT_MRT4530_102297C.1
US-10-437-963-2533
Percent Similarity:
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                                                                                 Pred. No.:
                                                                                                                                                                                                                                                         TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 203-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 3120
LENGTH: 792
TYPE: DNA
ORGANISM: Oryza sativa
FEATURE:
FORTHER THEOREMATTON: Oliver Transport File Company of the C
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
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Wu, Wei
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DB:
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Best Local Similarity:
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APPLICANT: Wu, Wei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Boukharov, Andrey A.
APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 49610
LENGTH: 568
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CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 63644
LENGTH: 297
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Publication No. US20040123343A1
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APPLICANT:
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APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
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ORGANISM: Oryza sativa
FEATURE:
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Cao, y
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Boukharov, Andrey A.
Barbazuk, Brad
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Query Match:
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Best Local Similarity:
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; OTHER INFORMATION: Clone ID: PAT_MRT4530_90693C.1
US-10-437-963-92187
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 92187
LENGTH: 571
TYPE: DNA
ORGANISM: Oryza sativa
                                                                                               GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas
APPLICANT: Kovalic, David
                                                                                                                                                 Sequence 64563, Application US/10437963
Publication No. US20040123343A1
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APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
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Zhou, Yihua
Cao, Yongwei
Wu, Wei
Boukharov, Andrey A.
Barbazuk, Brad
 Barbazuk, Brad
Li, Ping
                                Boukharov, Andrey A.
                                                   Wu, Wei
                                                                Zhou, Yihua
Cao, Yongwei
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US-10-282-122A-26385/c
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; OTHER INFORMATION: Clone ID: PAT_MRT4530_65695C.1
US-10-437-963-64563
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APPLICANT: Xu, Ĥ.

TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
FILE REFERENCE: ELITRA.034A
CURRENT APPLICATION NUMBER: US/10/282,122A
CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR PILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
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                     Remaining Prior Application data removed NUMBER OF SEQ ID NOS: 78614 SOFTWARE: PatentIn version 3.1
                                                                                        PRIOR FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/269,308
PRIOR FILING DATE: 2001-02-16
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DR FILLING DATE: 2.000-09-06
DR PELLING DATE: 2.000-09-07
DR FILLING DATE: 2.000-09-09
DR APPLICATION NUMBER: 60/242,578
DR FILLING DATE: 2.000-10-23
DR PELLING DATE: 2.000-10-23
DR PELLING DATE: 2.000-11-27
DR APPLICATION NUMBER: 60/253,625
DR FILLING DATE: 2.000-11-27
DR APPLICATION NUMBER: 60/257,931
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WARE: Pate
D NO 26385
                                                                                                                                                           APPLICATION NUMBER: 60/267,636
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Yamamoto, Robert
Forsyth, R.
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Zyskind, Judith
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Trawick, John
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Malone, Cheryl
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Percent Similarity:
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; Sequence 28450, Application US/10282122A
; Publication No. US20040029129A1
 Percent Similarity:
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US-10-282-122A-26385
                                                      Alignment Scores:
                                                                                            ; ORGANISM: Mycobacterium tuberculosis US-10-282-122A-28450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-857-841-4 (1-14) x US-10-282-122A-26385 (1-1603)
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SEQ ID NO 28450
LENGTH: 2625
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CURRENT APPLICATION UMBER: US/10/282,122A
CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR FILING DATE: 2000-05-23
                                                                                                                                                                                                        Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 78614
                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2000-12-22 PRIOR APPLICATION NUMBER: 60/267,636
                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/257,931
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PRIOR FILING DATE: 2000-09-06
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                                                                                                                                                                                                                                                                                        PRIOR FILING DATE:
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                                                                                                                                 TYPE: DNA
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Zyskind, Judith
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Malone, Cheryl
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Aro, Robert
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RESULT 15

(WS-10-437-963-86635/c)
(US-10-437-963)
(Publication No. US20040123343A1)
(Publication No. US20040123343A1)
(Publicant: La Rosa, Thomas J.
(APPLICANT: La Rosa, Thomas J.
(APPLICANT: Kovalic, David K.
(APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
(APPLICANT: Boukharov, Andrey A.
(APPLICANT: Boukharov, Andrey A.
(APPLICANT: Boukharov, Andrey A.
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Search completed: December 5, 2004, 09:47:10 Job time : 351 secs
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SEQ ID NO 86635
LENGTH: 969
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ORGANISM: Oryza sativa
FEATURE:
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-Q-(cgn2 1/USPTO_spool/US99857881/runat_03122004_160613_19119/app_query.fasta_1.199
-DB=GenEmbl -QFWT=fastap -SUFFIX.=clip2n.rge -MINWATCH=0.1 -LOOPCII-0 -LOOPEXT=0
-UNITS=bits -START=1 -END=-1 -MATRIX=clip0 -TRANS=human40.cdi -LIST=45
-DOCALIGN=200 -THR_SCORE=quality -THR_MIN=1 -ALIGN=15 -MODE=LOCAL -OUTFMT=pto
-NORN-ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=2000000000
-USER=US09857841_@CGN_1 _3731_@runat_03122004_160613_19119 -NCPU=6 -ICPU=3
-NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=60 -XGAPEXT=60 -FGAPOP=6
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# ALIGNMENTS

		COMMENT	JOURNAL	TITLE	REFERENCE		SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	TOTIS:	RESULT 1 BD261798
PC C12N15/09,A01H5/00,C07K14/415,C07K19/00,C12N5/10,C12N15/00, PC C12N5/00 CC Enhancement in protein production by higher plants using CC ubiquitin or cucumber mosaic virus coating protein peptide FH Key	PN JP 2002532098-A/2 PD 02-OCT-2002 PF 11-DEC-1998 JP 2000588378 PI RONG XIANG FANG, JUNG LIN WU, XIAO YING CHEN	INSTITUTE OF MOLECULAR AGROBIOLOGY OS Cucumber mosaic virus	or cucumber mosaic virus coating protein peptide Patent: JP 2002532098-A 2 02-OCT-2002;	<pre>Fang,R.X., Wu,J.L. and Chen,X.Y. Enhancement in protein production by higher plants using ubiquitin</pre>	1 (bases 1 to 53)	Viruses seRNA positive-strand viruses, no DNA stage; Bromoviridae;	Cucumber mosaic virus (cucumber mosaic cucumovirus)	JP 2002532098-A/2.	BD261798.1 GI:33071566	or cucumber mosaic virus coating protein peptide. BD261798	ent in protein production by higher plants u	BD261708 53 hp DNA linear DAT 17III2003	

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        1 (bases 1 to 130)
Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5849548-A 14 15-DEC-1998;
Location/Qualifiers
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Unclassified.
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:12305"
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                                         Unclassified.

1 (bases 1 to 130)
1 (bases 1, Brand, A., Perrimon, N. and Goodman, H.M. Haseloff, J., Brand, A., Perrimon, I. and Goodman, H.M. Cell ablation using trans-splicing ribozymes Patent: US 5641673-A 14 24-JUN-1997;
Location/Qualifiers
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I49964
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AR097446
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Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 6071730-A 14 06-JUN-2000;
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from patent US 6071730.
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AR031572
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AR065684
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Sequence 11
AR065684
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Sequence 11
AR031572
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Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5866384 A 11 02-FEB-1999;
Location/Qualifiers
                                                                                                      1 (bases 1 to 131)
Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5849548-A 11 15-DEC-1998;
Location/Qualifiers
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US-09-857-841-4 (1-14) x AR097443
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US-09-857-841-4 (1-14) x I49961 (1-131)
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Sequence 11 from patent US 5641673.
149961
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 6071730-A 11 06-UUN-2000;
Location/Qualifiers
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5641673-A 11 24-TUN-1997;
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AR031570
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Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.
Cell ablation using trans-splicing ribozymes
Patent: US 5866384-A 10 02-FEB-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 152)
Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5866384-A 9 02-FEB-1999,
Location/Qualifiers
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RESULT 13
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AR065683
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Haseloff, J., Brand, A., Perrimon, N. and Goodman, H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 5849548-A 10 15-DEC-1998;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 152)
Haseloff,J., Brand,A., Perrimon,N.
Cell ablation using trans-splicing
Patent: US 5849548-A 9 15-DEC-1998,
Location/Qualifiers
                                                                                             1 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
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Sequence 9 from patent
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AR097442
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Query Match:
Search completed: December Job time: 2542 secs
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Haseloff,J., Brand,A., Perrimon,N. and Goodman,H.M.
Cell ablation using trans-splicing ribozymes
Patent: US 6071730-A 10 06-UUN-2000;
Location/Qualifiers
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Sequence 10 from patent US 6071730.
AR097442 GI:12806172
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-Q=/cgn2 1/USPFO spool/US09857841/runat 03122004 160613 19108/app query.fasta 1.199
-DB=N Geneseq_23Sep04 -QFMT=fastap -SUFFIX=olip2n.rng -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=olip0 -TRANS=human40.cdi
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-NO MMAP -LARGEQUERY -NEG SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=60 -XGAPEXT=60 -FGAPOP=6
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### ALIGNMENTS

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                         11-DEC-1998;
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                                                                           22-JUN-2000.
                                                                                                                                                                                                        Nicotiana tabacum.
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                                                                                                    WO200036129-A1.
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                                                                                                                                       "Ubiquitin_monomer_fragment"
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02-DEC-1996
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                                          Boeshore ML,
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                                          Tricoli DM,
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                                        Reynolds JF,
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                                          Carney KJ;
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RESULT 3
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae, resistant to infection.
                                                                                                                                                                                                                                                                                                                                                                                             16-OCT-2003
02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1996-333993/33.
P-PSDB; AAR98895.
                                                                               WPI; 1996-333993/33.
P-PSDB; AAR98893.
                                                                                                                                                                                        07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                      Coat protein of the V27 strain of cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT34664;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 24; Fig 3; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      plants, partic. of the resistant to infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated cucumber mosaic virus coat protein DNA - used to produce plants, partic. of the family Cucurbitaceae or Solanaceae, which are
                                                                                                                  Boeshore ML,
                                                                                                                                                                                                                                      W09621018-A1
                                                                                                                                                                                                                                                                                                           Cucumber mosaic virus; strain V27.
                                                                                                                                                                                                                                                                                                                                              Coat protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT34664 standard;
                                                                                                                                                                                                               11-JUL-1996.
                                                                                                                                          (ASGR-) ASGROW SEED CO.
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CP) genes of the V27, V3 (CMV) were isolated. The
                                                                                                                   Mcmaster JR,
                                                                                                                                                                                                                                                                                                                                              CP; cucumber mosaic virus; CMV; V27; V33; V34; A35;
                                                                                                                                                                94US-00367789
                                                                                                                                                                                        95WO-US007234.
                                                                                                                                                                                                                                                                                      Location/Qualifiers
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Matches:
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Indels:
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                                                                                                                  Reynolds
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Claim 2; Fig 1;

80pp; English

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RESULT 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Percent Similarity:
                                             Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CNV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of CDNA using primers based known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CMV) were isolated. The CMV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based on known CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                  New isolated cucumber mosaic virus coat protein DNA - used to produce plants, partic. of the family Cucurbitaceae or Solanaceae, which are resistant to infection.
                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-333993/33.
P-PSDB; AAR93803.
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02-DEC-1996
  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             resistance;
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(first en
     BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of the A35 strain of cucumber mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mcmaster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEED CO
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Matches:
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RESULT 5
AAT34665
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Best Local Similarity:
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                                                        Alignment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Alignment Scores:
                                                                                                      Coat protein (CP) genes of the V27, V33, V34 and A35 strain of cucumber mosaic virus (CWV) were isolated. The CWV CP genes were isolated from infected plant tissue by PCR amplification of cDNA using primers based cknown CP sequences. The genes may be used for producing plants such as quash, cucumber, peppers and tomatoes which are resistant to CMV
                                                                                                                                                                                   New isolated cucumber mosaic virus coat protein DNA - used plants, partic. of the family Cucurbitaceae or Solanaceae, resistant to infection.
                                                                                                                                                                                                                                                                                                                                                                                    CDS
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                                                                           Sequence 773
                                                                                                                                                                Claim 13; Fig 2; 80pp; English.
                                                                                                                                                                                                                                                       Boeshore ML,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT34665 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                       WO9621018-A1
                                                                                                                                                                                                                                                                                                                                                                                                                 Cucumber mosaic virus; strain
                                                ŏ.
                                                                                               infection.
                                                                                                                                                                                                                                                                           (ASGR-) ASGROW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein of the V33 strain of cucumber mosaic virus.
                                                                                                                                                                                                                                     1996-333993/33.
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                                                                           BP; 175 A; 200 C;
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Matches:
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Gaps:
                                                                           G; 213 T;
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                                                                                                 field
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US-09-857-841-4 (1-14) x AAT34665 (1-773)

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Best Local Similarity:
Query Match:
DB:
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                                                  US-09-857-841-4 (1-14) x AAQ10461 (1-976)
                                                                                                       Percent Similarity:
                                                                                                                                       Alignment Scores
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 6
                                                                                                                                                                                The gene was isolated from a plasmid (pUC18) containing DNA complementary to RNA 3 of the virulent strain FNY, isolated in New York on infected melons. The sequence includes a leader sequence, coding region and 3' non-coding region. The leader sequence is the same length as that of the Japanese Y strain of CMV. Transgenic melon plantlets containing the sequence introduced via an Agrobacterium tumefaciens intermediate are cultured as shoots in special media. Transformed melon plants are resistant to CMV. See also AAQ10462. (Updated on 25-MAR-2003 to correct pA field.) (Updated on 25-MAR-2003 to correct PI field.) (Updated on 27-AMR-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                             Transgenic melon plantlets prodn. from transformed shoots -culture on specific medium for new transformed plants etc., resistant to cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                          Deboth M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Capsid protein gene of Cucumber Mosaic Virus strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-AUG-2003
25-MAR-2003
                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                         Claim 15; Page 17; 44pp; French.
                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1991-046027/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-AUG-1989;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-FEB-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EP412912-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CMV;
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                                                                                                                                                                                                                                                                                                                                                                                                                              (BIOC-) BIOCEM SA
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                                                                                                                               NO. :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 resistance; capsid protein; Cucumis melo;
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     MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg 14
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                                               US-09-857-841-4 (1-14)
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25-MAR-2003
16-APR-1991
                                                                                                                                                                                                       The gene was isolated from the virulent French strain 117F of CMV. Tomato plantlets at the 2-leaf stage were infected with CMV strain 117F. Is days post-infection, the virus was purified from the infected leaves and cDNA was synthesised from RNAs 1,2,3 and 4. After purification and selection by standard techniques, cDNA complementary to RNA 4 was found to encode the capsid protein. It was recloned into "Blue scribe" plasmids and sequenced. Transgenic melon plantlets containing the sequence introduced via an Agrobacterium tumefaciens intermediate are cultured as shoots in special media. The transformed melon plants are resistant to CMV. See also AAQ10461. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct OS
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 199
                                                                                                                                                                      Sequence 1007
                                                                                                                                                                                                                                                                                                                                                                                Transgenic melon plantlets prodn. from transformed shoots -culture on specific medium for new transformed plants etc., resistant to cucumber mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ10462 standard; DNA; 1007 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Deboth M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-AUG-1989;
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DB; AAR10653.
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            MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
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(revised)
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                                                                                                                                                                                                                                                                                                                                                                                                     New polyribozyme contg. several catalytic regions in complementary sequence - can inactivate gene for viral capsid protein, esp. for prepn. of new virus resistant transgenic plants, also DNA sequence encoding it.
 25-MAR-2003
21-MAY-1998
                                                                                                                                                                                                                                                                 Sequence 1007
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12-APR-1995
                                  AAT99545;
                                                        AAT99545 standard; cDNA; 1066
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                                                                                                                  MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
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e; inactivate;
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tivation; resistance; crop protection;
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US-09-857-841-4 (1-14) x AAT99545 (1-1066)

MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg

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                                                                                                                                This cDNA clone of cucumber mosaic virus (CMV) RNA-3 includes domain F of Cthe subgenomic promoter of viral RNA, the coat protein gene, its leader sequence, and the tRNA-like 3'-terminal region of RNA-3. It was prepared by amplifying a fragment of CMV cDNA-3 from clone pCR-CMVIRNA3 by PCR (see also AAT99548-49). The gene construct is introduced into a vector containing a promoter active in plant cells in antisense orientation relative to the promoter. A claimed recombinant vector comprises the plant promoter, the antisense gene construct and a terminator which is functional in the plant. In addition to CMV, viral RNA may also be used from tobacco mosaic virus and potato virus. Claimed transgenic plants have the antisense gene construct integrated into their genomes. They are resistant to viral infection. In particular, they are resistant to CMV. Within the antisense gene construct, the interfering activity of the antisense activity performed by the coat protein gene, allows production of plants having 100% resistance to CMV. (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                         Sequence 1066
                                                                                                                                                                                                                                                                                                                                                                                                             Preparing transgenic plants resistant to RNA virus infection - using anti:sense gene constructs containing the viral coat protein gene, e from cucumber mosaic virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               construct into the plant genome. The construct comprises: (a) an F domain of a subgenomic promoter of cucumber mosaic virus (CNV); (b) downstream from the subgenomic promoter, a leader sequence of a coat protein gene of CMV; (c) downstream from the leader sequence, a gene encoding a CMV coat protein; and (d) downstream from the gene, a 3'-terminal region of a CMV coat protein gene. The method is useful for producing plants which are resistant to infection by RNA based viruses. The gene construct gives higher levels of resistance compared to antisense constructs which are capable of complementing with different domains of genomic RNA of CMV. The present sequence represents the fragment of cDNA clone of RNA-3 of CMV. This forms the antisense construct of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to preparing transgenic plants resistant to virus induced infections that comprises integrating an antisense of the compression of the compress
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                                                                                    AAN81111;
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                                                                                                                                                                                                                                                                                                                                                                                                 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg 14
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25-FEB-1987;
18-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                             This DNA is produced on screening of a plasmid library and is used to transform plant cells which subsequently produce the coat protect cucumber mesaic virules (CMV) strain Y. This protein is not synthesised in patural plant cells. The resistance to CMV infect-ion, provided by the DNA, is shown in plant cells and redifferent- lated plant bodies. See also AAN81110. (Updated on 25-MAR-2003 to correct PR field.) (Updated calls and AN81110.
Cucumber mosaic virus; tomato mosaic virus; minus-strand RNA; virus resistance; disease resistance; transgenic plant; cross
                                                           25-MAR-2003
20-JUL-1995
                                    Cucumber mosaic virus RNA-3 chimera encoding RNase T1.
                                                                                                                       AAQ76107 standard;
                                                                                                                                                                                                                                                                                                                                            Sequence 1379 BP; 306 A; 342 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page ?; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA coding for the coat protein of cucumber mosaic virus strain Y - used for producing plants resistant to cucumber mosaic virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1988-236708/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cucumber mosaic
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12-NOV-1990
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87JP-00043443.
88JP-00035809.
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Matches:
Conservative:
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  cross protection;
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(SANO)
(SANO)
           virus resistance; disease resistance; transgenic plant; hypersensitive response; crop improvement; tomato; Lycopersicon esculentum; tobacco; Nicotiana tabacum; coa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA constructs for imparting pathogen, esp. virus, resistance to plants encodes RNA that interacts with viral RNA polymerase to generate an eliciting agent.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hypersensitive response;
Lycopersicon esculentum;
cell inhibitory protein;
                                                                                            25-MAR-2003
20-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                       A chimeric cucumber mosaic virus RNA-3 (given in AAQ76107) has the coat protein gene replaced by a gene encoding a cell inhibitory protein, RNase T1, having the sequence given in AAR67754. The construct elicits a minusenese RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus-resistance on a host plant, e.g. tobacco, tomato. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-JUN-1994;
                                              Cucumber mosaic virus; tomato mosaic virus; minus-strand RNA;
                                                                       Cucumber mosaic
                                                                                                                                                       AAQ76106
                                                                                                                                                                                                                                                                                                                                                                       Sequence 1696 BP; 387 A; 413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; Page 31-32;
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                                                                                                                                                                                                                                                                                                                                     No.
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OB; AAR67754.
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SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN
                                                                                                                                                       standard;
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                                                                     virus RNA-3 chimera encoding
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tobacco; Nicotiana tabacum; RNase Tl;
TOMV; CMV; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "RNase
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             coat
                                  cross protection;
           protein; CP;
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Percent Similarity:
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                                                                                                                                                                                                                                                                                                                                                                            encodes RN
eliciting
                                                                                                                                                                                                                                                                                             A chimeric cucumber mosaic virus RNA-3 (given in AAQ76106) codes for the coat protein (CP) of tomato mosaic virus (AAR67752) as well as its own CP (AAR67753). The construct elicits minus-sense RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus-resistance on a host plant, e.g. tobacco, tomato. (Updated on 25-MAR-2003 to correct PN field.)
                                                                    25-MAR-2003
20-JUL-1995
                                                                                                                                                                                                                                                                             Sequence 1860 BP; 448
                                                                                                                                                                                                                                                                                                                                                                                   DNA constructs for imparting pathogen, encodes RNA that interacts with viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SANO)
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                                                                                                                AAQ76108
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-DEC-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cucumber mosaic virus.
                                                                                                                                                                                                                                                   No.:
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DB; AAR67752, AAR67753.
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SANDOZ PATENT GMBH.
SANDOZ-ERFINDUNGEN VERW
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                                                                                                                standard;
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                                                                                                                                                                                                                                                                                                                                                           English.
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Matches:
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14
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Cucumber mosaic virus; tomato mosaic virus; minus-strand RNA; virus resistance; disease resistance; transgenic plant; cross hypersensitive response; crop improvement; tomato; Lycopersicon esculentum; tobacco; Nicotiana tabacum; P30; elic

elicitor;

ToMV

protection;

Cucumber mosaic virus RNA-3 chimera encoding ToMV

P30.

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RESULT 15
AAQ03641
ID AAQ03
XX CACum
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                                                                                                                       24-OCT-2003
27-AUG-2003
25-MAR-2003
07-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A chimeric cucumber mosaic virus RNA-3 (given in AAQ76108) has the coat protein gene replaced by a gene encoding an elicitor, ToMV P30, having the sequence given in AAR67755. The construct elicits a minus-sense RNA that interacts with the RNA-dependent RNA-polymerase of an invading virus, thus conferring virus-resistance on a host plant, e.g. tobacco, tomato. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA constructs for imparting pathogen, esp. virus, resistance to plants encodes RNA that interacts with viral RNA polymerase to generate an eliciting agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDS
                                          Cucumber mosaic virus;
                                                                    CMV-C;
                                                                                             Cucumber Mosaic Virus-C
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 2173 BP; 561 A; 438 C; 532 G; 642 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 5; Page 34-35; 50pp; English.
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P-PSDB; AAR67755.
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                                                                                                                                                                                                               AAQ03641 standard; DNA; 657
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(SANO ) SANDOZ PATENT GMBH.
(SANO ) SANDOZ-ERFINDUNGEN VERW GES MBH
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                                                                                                                                                                                                                                                                            Coat protein gene of cucumber mosaic virus strain \mathtt{WL} - cloned transformed plants which are resistant to CMV viral infection.
                                                                                                                                                                                   Sequence 657 BP; 157 A; 176
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-DB=EST -QFMT=fastap -SUFFIX=01jp2n.rst -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS=bits -START=1 -END=-1 -MATRIX=oligo -TRANS-human40.cdi -LIST=45
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-DEV TIMEOUT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=60 -FGAPOP=6
-PGAPEXT=7 -YGAPOP=60 -YGAPEXT=60 -DELDP=6 -DELEXT=7
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Contact: Doil Choi
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pacs2-164_2261, genomic survey sequence.
BZ560224
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Box 352145, Seattle, WA 98105-2145,
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/clone="pacs2-164 2261"
/clone lib="pacs2-164"
/clone lib="pacs2-164"
/note="clinical isolate 2-164 Whole genomic shotgun library."
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/clone_lib="KS08"
/note="Vector: pBluescript SK(-)"
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Mammalia; buchler, Akimura, T., Arai, A., Arakawa, T., 1 (Dases 1 to 172)

Aizawa, K., Akahira, S., Akimura, T., Hiraoka, T., Hirozane, T., Carninci, P., Hanagaki, T., Hayatsu, N., Hiraoka, T., Hirozane, T., Hodoyama, Y., Imotani, K., Ishii, Y., Itoh, M., Izawa, M., Kawai, J., Kojima, Y., Konno, H., Kusakabe, M., Matsuyama, T., Miyazaki, A., Nomazaki, R., Okazaki, Y., Nakamura, M., Nishi, K., Nomura, K., Numazaki, R., Okazaki, Y., Okido, T., Owa, C., Sakai, C., Sakai, K., Sasaki, D., Sato, K., Shibata, K., Shibata, Y., Shinagawa, A., Shiraki, T., Sogabe, Y., Suzuki, H., Tagawa, A., Takahashi, F., Tanaka, T., Toya, T., Suzuki, H., Tagawa, A., Takahashi, F., Tanaka, T., Yoshiki, A., Washahiki, A., Yamamura, T., Yasunishi, A., Yoshida, K., Yoshiki, A.,
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Spencer, D.H., Raymo
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msh2_191.y1 msh Pseudomonas
genomic survey sequence.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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/note="Environmental isolate. Whole genomic shotgun
library."
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/mol_type="genomic DNA"
/strain="MSH"
/db_xref="taxon:287"
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Thermostabilization and thermoactivation of thermolabile enzymes by trehalose and its application for the synthesis of full length cDNA. Proc. Natl. Acad. Sci. U.S.A. 95 (2), 520-524 (1998)
Itoh,M., Kitsunai,T., Akiyama,J., Shibata,K., Izawa,M., Kawai,J., Tomaru,Y., Carninci,P., Shibata,Y., Ozawa,Y., Muramatsu,M., Okazaki,Y. and Hayashizaki,Y.
Automated filtration-based high-throughput plasmid preparation system. Genome Res. 9 (5), 463-470 (1999)
Carninci,P. and Hayashizaki,Y.
High-efficiency full-length cDNA cloning. Methods Enzymol. 303, 19-44 (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Muramatsu, M. and Hayashizaki, Y.
RIKEN Mouse ESTs (Aizawa, K. et al. 2000)
Unpublished (2000)
Contact: Yoshihide Hayashizaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Please visit our web site (http://genome.rtc.riken.go.jp) for further details.
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                                        ThrSerAlaGlyArgAsnArgArg 14
                                                                                                                                                                                                                                                                                                                                       /clone lib="RIKEN full-length enriched, 11 days pregnant adult female ovary and uterus"
/note="Site 1: Sall; Site 2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN, Division of Experimental Animal Research in Riken
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /tissue_type="ovary and uterus"
/dev_stage="11 days pregnant, adult"
/lab_host="DH10B"
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(mol_type="mRNA"
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IMAGE:972821 5', mRNA sequence.
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This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=RC3&t2=RC3-ST0186-
181099-012-c02&t3=1999-10-18&t4=1)
Seq.primer: puc 18 forward
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 402)
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010,
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The FAPESP/LICR Human Cancer Genome Project
Unpublished (1999)
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RC3-GT0186-181099-012-c02 ST0186 Homo sapiens cDNA,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="mRNA"
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/dev_stage="Adult"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        269
                                                                                                                                                                                                                                                                                                                                                                                               AQ612674 536 bp DNA linear GSS 15-JUN
HS_5107_B2_E12_T7A RPCI-11 Human Male BAC Library Homo sapiens
genomic_clone Plate=683 Col=24 Row=J, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WashU-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubu
Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M.,
Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B.,
Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Waterston, R.
The WashU-HHMI Mouse EST Project
                                                                                                                               Sequence-tagged connectors: A sequence approach to mapping
                                                                                                                                                                                                                                                                                                                 HOMO
                                                                                                                                                                                                                                                                                                                                                        AQ612674
AQ612674.1 GI:5073950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Marra M/Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1996
                                                                                                                                                                      Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 536)
Mahairas,G.G., Wallace,J.C., Smith,K., Swartzell,S.,
Keller,A., Shaker,R., Furlong,J., Young,J., Zhao,S.,
                                                                                                           scanning the
                                                                                                                                                                                                                                                                    Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MGI:553549
                                               10449764
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Location/Qualifiers
act: Mahairas GG, Wallace JC,
Throughput Sequencing Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 314 286 1800
314 286 1810
                                                                                                                                                                                                                                                                                           sapiens
                                                                                                                                                                                                                                                                                                           sapiens (human)
                                                                                      Natl. Acad. Sci. U.S.A. 96 (17),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /dev_stage="M30 CD4+ cells"
/lab host="SOLR (kanamycin resistant)" |
/lab host="SOLR (kanamycin resistant)" |
/clone lib="Stratagene mouse Tcell 97311" |
/note="Organ: blood; Vector: pBluescript SK-; Site_1:
EcoRI; Site_2: XhOI; Cloned uniddrectionally. Primer:
Oligo dT. M30 CD4+ cells. Average insert size: 1.0 kb;
Uni-ZAP XR Vector; -5' adaptor sequence: 5' GAATTCGGCACGAG
3' ~3' adaptor sequence: 5' CTCGAGTTTTTTTTTTTTTTTT 3'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mol_type="mRNA"
/db_xref="taxon:10090"
/clone="IMAGE:972821"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tissue_type="Tcell"
                                                                                                                                                                                                                                             Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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8.00
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                                                                                                           human genome
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                               Holzman, T., Adams, M.D.
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AUTHORS
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ORGANISM
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LOCUS
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        486 GAGAGCACATCTGCCGGCCGCAAT 463
                                                                                    Contact: Kirkness EF
The Institute for Genomic Research
Department of Eukaryotic Genomics,
Rockville, MD 20850, USA
Tel: 301-338-0200
                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Carnivora; Fissipedia; Canidae; Canis. 1 (bases 1 to 589)
Kirkness, E.F., Bafna, V., Halpern, A.L., Levy, S., Remington, K., Rusch, D.B., Delcher, A.L., Pop, M., Wang, W., Fraser, C.M. and Venter, J.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: jwallace@u.washington.edu
Clones are derived from the human BAC library RPCI-11. For BAC
library availability, please contact Pieter de Jong
(pieter@dejong.med.buffalo.edu). Clones may be purchased from
BACPAC Resources (http://bacpac.med.buffalo.edu/ordering_bac.htm)
or from Resear h Genetics (info@resgen.com). BAC end Web Server:
http://www.htsc.washington.edu
Plate: 683 row: J column: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genomic survey
CE588296
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tigr-gss-dog-17000366361600 Dog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 GluSerThrSerAlaGlyArgAsn 12
                           Class: shotgun.
                                           Email: ekirknes@tigr.org
                                                                                                                                                                                                                                             The dog genome: survey sequencing and comparative analysis science 301 (5641), 1898-1903 (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                             Canis familiaris
                                                                                                                                                                                                                                                                                                                                                                                                                                               Canis familiaris (dog)
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Class: BAC ends
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401 Queen Anne Avenue North, Seattle, WA 98109, USA
Tel: (206) 616-318
Fax: (206) 616-3887
                                                                                                                                                                                                      14512627
                                                                      301-838-0208
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone lib="RPCI-11 Human Male BAC Library"
/note="Vector: pBACe3.6; Site 1: EcoRI; Site 2: EcoRI;
Male blood DNA was isolated from one randomly chosen do
and partially dispested with a combination of EcoRI and
EcoRI Methylase. Size selected DNA was cloned into the
pBACe3.6 vector at EcoRI sites"
    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="genomic DNA"
/db_xref="taxon:9606"
/clone="Plate=683 Col=24 Row=J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /sex="male"
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Mismatches:
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Library Canis familiaris
                                                                                                                                      9712 Medical Center Drive,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   00008536
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REFERENCE
AUTHORS
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CF568862
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ORGANISM
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Best Local Similarity:
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DB:
                                              Percent Similarity:
Best Local Similarity:
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                                                                                                                                        Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Jiang Lu, FAMU, Wayne Hunter, Viticulture Centre, FAMU, Tallahassee, Florida A&M University
Tallahassee, FL 32317, USA
Tel: (850) 412-7393
Fax: (850) 561-2617, (772) 462-5898
Email: jiang.lu@famu.edu,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hunter,W.B., Dang,P.M., Chaparro,J.X., Lu,J. and Genes expressed in Vitis shuttleworthii L Unpublished (2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; Vitaceae; Vitis.
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FAMU UDDA FP 00084 Vitis shuttleworthii L.,
shuttleworthii cDNA 5', mRNA sequence.
CF568862
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer:
                                                                                                                                                                                                                      /sex="Mixed population"
/tissue type="Entire tendril, leaves, bud, flowers"
/dev stage="At blooming"
/lab_host="XLI-Blue"
/clone lib="Vitis shuttleworthii L., grape"
/note="Vector: pBluescript II SK+; Site 1: EcoRI; Site 2:
XhoI; A high quality EST with at least 100 contiguous
bases at Trace Tuner score of 20 or better. Construction
by PM Dang, USDA, ARS, U.S. Horticultural Research Lab,
"Times" USDA, ARS, U.S. Horticultural Research Lab,
                                                                                                                                                                                                   by PM Dang, USDA, AND
Ft.Pierce, FL, USA."
                                                                                                                                                                                                                                                                                                                                                                                                                                          /mol_type="mRNA"
/db_xref="taxon:246827"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peripheral blood"
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/note="Site_1: BstXI; Libraries were prepared from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mol_type="genomic DNA"
/strain="Standard Poodle"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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AZ419370/c
Alignment Scores: Pred. No.:
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: C
Plate: 0195 row: P column: 18
Seq primer: CACACAGGAAACAGCTATGACC
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High quality sequence stop: 679.
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
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                                                                                was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNP42 (gi |4732114|gb|AF7129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /Clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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  311
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Insert Length: 10000 Std Error:
Plate: 0233 row: J column: 19
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University of Utah Genome Center
University of Utah
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686 bp DNA linear GSS 27-APR-20
2M0233J19R Mouse 10kb plasmid UUGC2M library Mus musculus genomic
clone UUGC2M0233J19 R, genomic survey sequence.
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Seq primer: CACACAGGAAACAGCTATGACC
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Mammalia; Eutheria; Rodentia;
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801 585 7177
                       /lab host="E. coli strain XL10-Gold, T1-resistant, F-"
/(lab host="E. coli strain XL10-Gold, T1-resistant, F-"
/(clone_lib="Mouse_10kb plasmid_UUGC2M_library"
/note="Vector: PMD42nv; Purified_genomic_DNA_from M.
musculus CS7BL/6J (female) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
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Percent Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The Institute for Genomic Research 9712 Medical Center Dr., Rockville, Tel: 301 838 0200 Fax: 301 838 0208 Email: szhao@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AZ741218 816 bp DNA line RPCI-24-145M10.TJ RPCI-24 Mus musculus genomic RPCI-24-145M10, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Clones are derived from the mouse BAC library RPCI-24. For BAC library availability, please contact Pieter de Jong (pdejong@mail.cho.org). Clones may be purchased from BACPAC Resources (http://www.chori.org/bacpac/orderingframe.htm). BAC page: http://www.tigr.org/cdb/bac_ends/mouse/bac_end_intro.html Plate: 145 row: M column: 10
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Other_GSSs: RPCI-24-145M10.TV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seq primer: SP6
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Department of Eukaryotic Genomics
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Zhao,S., Nierman,W., Malek,J., Shatsman,S., Akinret,B., Levins
Tsegaye.G., Geer,K., Krol,M., Shvartsbeyn,A., Gebregeorgis,E.,
Russell,D., de Jong,P. and Fraser,C.M.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                       /cell_type="Spleen/Brain"
/clone_lib="RRCI-24"
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
RPCI-24 Mouse BAC Library produced by Pieter de Jong. The
library was cloned in the pTARBAC1 cloning vector at the
BamH1 sites using MboI partially digested male C57BL/6J
DNA."
                                                                                                                                                                                                                                                                                                                                               /organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
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/clone="RPCI-24-145M10"
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 BZ603235 BZ603215 Human MCF7 breast cancer cell line library (MCF7_1) Homo sapiens genomic clone MCF7_1-19021, genomic survey sequence.

BZ603235
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Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Gilbert Smith, Ph.D.

cDNA Library Preparation: Life Technologies, Inc.

cDNA Library Arrayed by: The I.M.A.G.B. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:

http://image.lnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 823)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
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603378972F1 NCI_CGAP_Mam2 Mus musculus cDNA clone IMAGE:5391617 5',
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/note="Organ: mammary; Vector: pCMV-SPORT6; Site_1: Sall;
Site_2: Noti; Cloned unidirectionally. Primer: Oligo dT.
Library constructed by Life_Technologies. Investigator
                                                                                                                                                                                                                                                                                                                                                                 providing samples: Gilbert
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/dev_stage="5 months"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="mRNA"
/strain="FVB/N-3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    clone="IMAGE:5391617"
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Unpublished Other_GSSs:
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1 (bases 1 to 839)

Volik,S., Zhao,S., Chin,K., Brebner,J.H., Herndon,D.R., Tao,Q., Kowbel,D., Huang,G., Lapuk,A., Kuo,W.-L., Magrane,G., de Jong,P., Gray,J.W. and Collins,C.

End-sequence profiling: Sequence-based analysis of aberrant genomes proc. Natl. Acad. Sci. U.S.A. 100 (13), 7696-7701 (2003)
                                                                             1 (bases 1 to 889)
Whitelaw, C.A., Quackenbush, J., Van Aken, S., Utterback, T.,
Resnick, A., Fraser, C.M., Yuan, Y., San Miguel, P., Ma, J. and
                                                                                                                                                          Zea mays
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD
clade; Panicoideae; Andropogoneae; Zea.
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CG137878.1 GI:34028661
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Tel: 415 502 7066
Fax: 415 502 5665
                                             Maize Genomics Consortium
                                                                                                                                                                                                                                                                                                                                                                       PUICG49TD ZM_0.6_1.0_KB Zea mays genomic clone ZMMBTa0550J01,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: svolik@cc.ucsf.edu
This clone is available fi
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/mol type="genomic DNA"
/db_xref="taxon:9606"
/clone="MCF7 1-19021"
/sex="female"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Vector: pBCBAC1; Site_1: HindIII; This library was constructed from MCF7 breast cancer cell line by Amplicon Express (http://www.genomex.com) using their standard procedure."
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Contact: Cathy Whitelaw
TIGR
9712 Medical Center Drive, Rockville, MD 20850, USA
Tel: 301-838-5943
Fax: 301-838-5943
Fax: 301-838-50208
Email: whitelaw@tigr.org
Seq primer: TF
Class: sheared ends. lifers
1. 089
1. 089
1. 089
1. 089

FEATURES

SOURCE

Class: sheared ends. lifers
1. 0001_type="genomic DNA"
/strain="R73"
/clone="ZMMTPA0550J01"
/
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No.
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Maximum DB
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-Q-/cgn2_1/USPTO_spool/US09857841/runat 03122004 160614 19150/app_query.fasta 1.199
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-FGAPEXT=7 -YGAPOP=60 -YGAPEXT=60 -DELOP=6 -DELEXT=7
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Perfect score:
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seq length: 2000000000
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Match
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2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/BTUS_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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           US-08-090-193-14
US-08-488-031-14
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US-08-488-027-14
US-08-090-192-14
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US-08-482-658-14
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US-08-090-193-14
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equence 12,	ence 12,	equence 10,	equence 10,	equence 9,	equence 10,	equence 9,	equence 10,	equence 9,	equence 10,	equence 9,	equence 10,	equence 9,	equence 10,	equence 9,	equence 10,	equence 9,	equence 10,	equence 9,	equence 10,	equence 9,	equence 10,	equence 9,	equence 11,	equence 11,	equence 11,	equence 11,	equence 11,	equence 11,	equence 11,	equence 11,	quence 11,	equence
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### ALIGNMENTS

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Sequence 14, Application US/08090193
Patent No. 5641673
GENERAL INFORMATION:
APPLICANT: Haseloff, James
APPLICANT: Brand, Andrea
APPLICANT: Brand, Andrea
APPLICANT: Goodman, No. 5641673bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox
STREET: 1100 New York Avenue, Suite 600
CITY: Washington
STATE: D.C.
COUNTRY: Washington
STATE: D.C.
ZIP: 20005
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/090,193
FILING DATE: 23-DEC-1993
CLASSIFICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JAN-1992
PRIOR APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
REGISTRATION NUMBER: 33,851
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Percent Similarity:
Best Local Similarity:
Query Match:
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Patent No. 58
                                                                                                                                                                                                                                                                                 SOFTWARE: PATENTIN Release #1.0, Ve.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,031
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,193
FILING DATE: 23-DEC-1993
PRIOR APPLICATION NUMBER: PCT/US92/00277
APPLICATION NUMBER: PCT/US92/00277
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TELEPHONE: (202) 371-2600
TELEPAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                              TELEFAX: (202) 371-254
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                            NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080002/MAC/LBB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
TELEPAX: (202) 371-2540
                                                                                                                                                                                           FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy
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TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
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TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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STRANDEDNESS:
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1100 New York Avenue,
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Matches:
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Indels:
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Percent Similarity:
Best Local Similarity:
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Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                       Alignment Scores:
                                                                                                                            US-08-486-569-14
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                                                                                                                                                                                                                                                                                 APPLICATION UNMER: 08/090,193

PILING DATE: 23-DEC-1993

PRIOR APPLICATION DATA:
APPLICATION UNWBER: PCT/US92/00277

FILING DATE: 16-JAN-1992

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,330

FILING DATE: 17-JAN-1991

ATTORNEY/AGENT INFORMATION:
NAME: Bugaisky, Lawrence B.
REGIGSTRATION NUMBER: 35,086

REFERENCE/DOCKET NUMBER: 0609.3080006/MAC/LBB

TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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GENERAL INFORMATION:
APPLICANT: Haselo
APPLICANT: Brand,
                                                                                                                                                                                                            TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/486,569
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Perrimon, No. 5863774bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
                                                                        No.:
                                                                                                                                                            LENGTH: 130 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 1100 New CITY: Washington
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1100 New York Avenue, N.W., Suite 600
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Mismatches:
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RESULT 5
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                                                                                                                                                                                                          US-09-857-841-4 (1-14) x US-08-488-027-14 (1-130)
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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FILING DATE: 23-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAW-1991
ATTORNEY/AGENT INFORMATION:
NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080003/MAC/LBB
TELECOMMUNICATION INFORMATION:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,027
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
ADDITION TO THE THE TO THE
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SOFTWARE: Patentin Balana "Topera"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
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T: Perrimon, No. 5866384bert
T: Goodman, Howard M.
INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
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                                                                                                                MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg
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Matches:
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Best Local Similarity:
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                                                                                                                                                                 Sequence 14, Application US/08482663
Patent No. 5882907
GENERAL INFORMATION:
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Patent No. 5874414
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APPLICANT: Haseloff, James
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Trans-Splicing Ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Cimbala, Michele A.
REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0609.3030001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox
STREET: 1100 New York Avenue, N.W., Suite 600
                                                   APPLICANT: Perrimon, No. 5882907bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 07/642,333 FILING DATE: 17-JAN-1991 ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: PCT/
FILING DATE: 16-JAN-1992
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STRANDEDNESS: both
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                                                                                                          Brand, Andrea
Perrimon, No. 5882907bert
                                                                                                                                                    Haseloff, James
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11-NOV-1993
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Matches:
Conservative:
Mismatches:
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COUNTRY:

U.S.A.

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Floppy disk

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US-08-482-658-14
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DB:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                            Sequence 14, Application US/08482658
Patent No. 6010904
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 07/64
PILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Bugalsky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/482,663
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,193
FILING DATE: 23-DEC-1993
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                  APPLICANT: Haseloff, James
APPLICANT: Brand, Andrea
APPLICANT: Perrimon, No. 6010904bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
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                                                                                                COUNTRY: U.
ZIP: 20005
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STRANDEDNESS: both
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                            ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                53 ATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGA
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D.C.
U.S.A.
                                                                                                                                                       Washington
                                                                                                                                                                       E: Sterne, Kessler, Goldstein & Fox 1100 New York Avenue, N.W., Suite 600
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PatentIn Release #1.0, Version #1.25
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Matches:
Conservative:
Mismatches:
Indels:
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Percent Similarity:
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Query Match:
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US-08-470-349-14
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/090,192
FILING DATE: 11-NOV-1993
APPLICATION NUMBER: PCT/ US 92/00278
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,333
                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Haseloff, James
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 51
CORRESPONDENCE ADDRESS:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
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APPLICATION NUMBER:
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TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                             STREET: 1100 New
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080008/MAC/LBB
                                                                                                                                                  APPLICATION NUMBER: FILING DATE: 06-JUI
                                                                                                                                                                                                                                                                                                 COUNTRY: U
ZIP: 20005
                                                                                                                                                                                                                                                                                                                                   STATE: D.C.
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(202) 371-2600
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CURRENT APPLICATION NUMBER: US/08/475,610
FILING DATE: 07-JUN-1995
PRIOR APPLICATION NUMBER: 08/090,193
FILING DATE: 23-DEC-1993
PRIOR APPLICATION NUMBER: 07/US92/00277
FILING DATE: 23-DEC-1993
PRIOR APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JAN-1992
PRIOR APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080004/MAC/LBB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2540
INFORMATION FOR SEQ ID NO: 14:
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Best Local Similarity:
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INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Cimbala, Michele A.
REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0609.3030001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371.2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox
STREET: 1100 New York Avenue, N.W., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Perrimon, No. 6071730bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
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TYPE: nucleic acid
STRANDEDNESS: both
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.25
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Matches:
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                                             Percent Similarity:
Best Local Similarity:
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PCT-US92-00277-14
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Best Local Similarity:
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                                                                                                    Pred. No.:
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LENGTH: 130 base pairs
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                                                                                                                                                                                                           TYPE: NUCLEIC ACID STRANDEDNESS: both
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TOPOLOGY: li
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ZIP: 20036
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US-09-857-841-4 (1-14) x US-08-475-610-14 (1-130)
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                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,
PILING DATE: I7-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jorge A
REGISTRATION NUMBER: 29/021
REFERENCE/DOCKET NUMBER: 0609
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)833-7533
                                                                                                                                                                                                                          TELEPHONE: (202)833-753
TELEFAX: (202)833-8716
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 130 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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APPLICANT: Perrimon, Norbert
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: linear
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Matches:
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Indels:
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Matches:
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US-09-857-841-4 (1-14) x PCT-US92-00277-14 (1-130)

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RESULT 12
US-08-090-193-11
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Best Local Similarity:
Query Match:
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                                                    Sequence 11, Applicat:
Patent No. 5641673
GENERAL INFORMATION:
APPLICANT: Haselo
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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FILING DATE: 19320116
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
FILING DATE: 17-CAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: GOLdstein, Jorge A
REGISTRATION NUMBER: 29,02:
 APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)833-7533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 130 bases
TYPE: NUCLEIC ACID
STRANDEDNESS: Single
TOPOLOGY: Linear
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                                                                                                       Application US/08090193
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Brand, Andrea
Perrimon, No. 5641673bert
Goodman, Howard M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
                                                      Haseloff, James
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Matches:
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Indels:
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Percent Similarity:
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Query Match:
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                                                                                                                                                                                                                                                             Patent No. 5849548
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,330
FILING DATE: 17-JAN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/090,193
FILING DATE: 23-DEC-1993
COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                  APPLICANT: Perrimon, No. 5849548bert
APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 131 base pairs
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CLASSIFICATION: 514
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                            APPLICANT: Haseloff, James APPLICANT: Brand, Andrea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                    STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
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REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0609.3080001
                                    COUNTRY: U
ZIP: 20005
                                                                                   CITY: Washington
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                                                                                                                       ADDRESSEE:
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                                                                                                                                                                                                     Brand, Andrea
Perrimon, No. 5849548bert
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Matches:
Conservative:
Mismatches:
Indels:
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CURRENT APPLICATION DATA:

PatentIn Release #1.0, Version #1.25

COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS

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COMPUTER READABLE FORM:

MEDIUM TYBE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/486,569
FILING DATE: 07-JUN-1995
PRIOR APPLICATION NUMBER: 08/090,193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 14
US-08-486-569-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 11, Application US/08486569 Patent No. 5863774
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INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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APPLICATION NUMBER: PCT/US92/00277
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/642,330
                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox
STREET: 1100 New York Avenue, N.W., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 060
TELECOMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/488, FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA: 08/090,193
APPLICATION NUMBER: 08/090,193
FILING DATE: 23-DEC-1993
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ATTORNEY/AGENT INFORMATION:
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STRANDEDNESS: both
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
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T: Goodman, Howard M.
T: NVENTION: Cell Ablation Using Trans-Splicing Ribozymes
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Matches:
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Best Local Similarity:
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US-08-488-027-11
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APPLICANT: Haselof
APPLICANT: Brand,
APPLICANT: Perrimo
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INFORMATION FOR SEQ ID NO:
                                                                                          PRIOR AFFLICTION NUMBER: 08/090,123
PRILING DATE: 23-DEC-1993
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
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PRIOR APPLICATION NUMBER: 08/090,193
                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FOR COMPATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 131 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3080006/MAC/LBB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2540
TELEFAX: (202) 371-2540
ATTORNEY/AGENT INFORMATION: NAME: Bugaisky, Lawrence
                                             APPLICATION NUMBER: PCT/I
FILING DATE: 16-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Goodman, Howard M.
TITLE OF INVENTION: Cell Ablation Using Trans-Splicing Ribozymes
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                                  FILING DATE:
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FILING DATE: 16-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
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Perrimon, No.
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Matches:
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REFERENCE/DOCKET NUMBER: 0609.3080003/MAC/LBB
TELLECHMOUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
TELEPAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 131 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
TTYPE: nucleic acid
                                                                                                   Fred. No.:

Score:
14.00
Percent Similarity: 100.00%
Best Local Similarity: 100.00%
Query Match:

DB:

3.34e-06
                                                                                                                                                                        Alignment Scores:
Pred. No.:
Score:
                                                                 US-09-857-841-4 (1-14) x US-08-488-027-11 (1-131)
Length:
Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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Search completed: December 5, 2004, 11:11:50 Job time : 68 secs

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Database :
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Maximum DB seq length: 200000000
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Ygapop 60.0 ,
Fgapop 6.0 ,
Delop 6.0 ,
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3: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*

5: /cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
                                                                                          US-09-857-841-4
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/ Ggn2_6/ptodata/2/pubpna/US09E_PUBCOMB.seq:*
/ Ggn2_6/ptodata/2/pubpna/US09E_PUBCOMB.seq:*
/ Ggn2_6/ptodata/2/pubpna/US09C_PUBCOMB.seq:*
/ Ggn2_6/ptodata/2/pubpna/US10A_PUBCOMB.seq:*
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Ygapext 60.0
Fgapext 7.0
Delext 7.0
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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US-1U-425-115-138/US	10 10 10 10 10 10 10 10 10 10 10 10 10 1	S-10-282-122A-752	A-26	-10-343-710-	S-10-369-493-4419	8-10-369-493-357	S-10-227-577-455	-10-091-504-	20 20 20 20 20 20 20 20 20 20 20 20 20 2	200 101 101 00 AND 00 A	10-186-761-01	S-10-437-963-6738	8-10-369-493-3	5-09-822-846-46	8-846-1082		-10-424-599-1109	US-10-431-096-37	-10-126-103-3	-10-424-599-4	S-10-369-493-3	US-10-425-115-61926	-10-156-761-4	-10-425-115-4	-10-156-761-7	S-10-425-115-596	S-10-029-386-	S-10-424-599-317	S-10-029-386-117	-10-027-632-23445	-10-027-632-2344	S-10-425-115-17448	-09-864-761-15	09-998-598-1555	-09-918-995-34	S-10-425-115-4648	09-960-352-7184	S-10-424-599-946	S-10-425-115-66023	S-10-424-599-140	S-10-437-963-	S-10-011-033-	S-10-011-033-3	S-10-011-033-	S-10-011-033-	US-10-011-033-5	ID
ednence ra	A 13070E	e 7523,	e 2638	e 78	equence 4415	equence 35//0,	equence 455,	quence 4	quence 400, s	100	777	nce 67382	32000	464,	quence 10	nce 1391	nce 1109	nce 37	nce 37,	4268	3773	61	4748, A	42949,	nce 7305, A	ı n	nce 1	equence 31741	equence 1	equence 23445	equence	Sequence 174482	equence 15219,	quence 1555, Ap	equence 34460,	Sequence 46484,	quence 7184, Ap	e 94618,	equence 66023,	e 140904	equence 3120, A	equence 9,	equence 3,	equence	equence 1,	Seguence 5. Appli	cription

### ALIGNMENTS

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US-10-011-033-5
Sequence 5, Application US/10011033
Publication No. US20020124286A1

GENERAL INFORMATION:
APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains of Cucumber Mosaic Virus
CORRESPONDENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
CITY: Chicago
STATE: Illinois
COUNTRY: US.A.
ZIP: 60089
COMPUTER READABLE FORM:
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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US-10-011-033-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 5:

SEQUENCE CHARACTERISTICS:
LENGTH: 771 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ORIGINAL SOURCE:
ORGANISN: Cucumber mosaic virus
                                                                                                                                                                                                                                                                       Sequence 1, Application US/10011033 Publication No. US20020124286A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-NO. US20020124286A1-2001
CLASSIFICATION: <UNknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMUNICATION INFORMATION:
TELEPHONE: 312-616-5460
NEORGATION 528-660
TELEPAX: 312-616-5460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8
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                                                                                                                                   APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
Carney, Kim J
CITLE OF INVENTION: Plante Resistant to C Strains
Cucumber Mosaic Virus
                                CORRESPONDENCE ADDRESS:

ADDRESSEE: Rockey, Milnamow & Katz, Ltd.

STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /standard_name= "COAT PROTEIN" SEQUENCE DESCRIPTION: SEQ ID NO: 5:
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                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                   1 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg 14
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LOCATION: 3..660
OTHER INCOMMATION: /codon_start= 3
/function= "ENCAPSIDATES VĪRUS RNA"
/product= "COAT PROTEIN"
/gene= "CP"
STATE: Illinois COUNTRY: U.S.A.
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Matches:
Conservative:
Mismatches:
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Gaps:
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COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA PC-DOS/MS-DOS
SOFTWARE: PATENTIA PC-DOS/MS-DOS
SOFTWARE: PATENTIA PLANTION DATA:
APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION: <UNKNOWN>
PRIOR APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
TELEPHONE: 312-616-5400
TELEPHONE: 312-616-5400
TELEPAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-011-033-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 3
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                             Sequence 14, Application US/10011033
Publication No. US20020124286A1
                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 772 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
ZIP: 60089

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Cucumber Mosaic Virus
INDIVIDUAL ISOLATE: V-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                      TITLE OF INVENTION: Plants Resistant to C Strains Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Boeshore, Maury L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 ATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGG
                                                                                               CITY: Chicago
STATE: Illinois
COUNTRY: U.S.A.
                                                                                                                                                             ADDRESSEE: Rockey, Milnamow & Katz, Ltd. STREET: 2 Prudential Plaza, 180 N. Stetson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: CDS
LOCATION: 3..660
                                                                                                                                                                                                                                                                               Reynolds, John Carney, Kim J
                                                                                                                                                                                                                                                                                                                   McMaster, J. Russ
Tricoli, David M
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Matches:
Conservative:
Mismatches:
Indels:
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US-10-011-033-3
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 3, Application US/10011033 Publication No. US20020124286A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/875,233
PILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5460
TELEPHONE: 312-616-5460
ZIP: 60089

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
BPILING DATE: 13-No. US20020124286A1-2001
                                                                                                                                                                                                                                                                                                     APPLICANT: Bosshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 772 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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PILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 15 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HYPOTHETICAL: NO ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 MetAspLysSerGluSerThrSerAlaGlyArgAsnArgArg 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATGGACAAATCTGAATCAACCAGTGCTGGTCGTAACCGTCGA
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Cucumber Mosaic Virus INDIVIDUAL ISOLATE: A35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                           STREET: 2 Pruder
CITY: Chicago
STATE: Illinois
                                                                                                                                                                           COUNTRY: U.S.A.
                                                                                                                                                                                                                            ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
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Matches:
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Indels:
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9, Application US/10011033 Publication No. US20020124286A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: 312-616-5400
TELEPAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 3:
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 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION NUMBER: US/10/011,033
APPLICATION UMBER: US/20020124286A1-2001
FILING DATE: 13-NO. US20020124286A1-2001
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: CINA
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Reynolds, John F
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 792 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
ATTORNEY/AGENT IMPORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE DESCRIPTION: SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 MetAspLysSerGluSerThrSerAlaGlyArgAspArgArg 14
                                                                                                                                                                                                                                                                     CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN: v-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                  ZIP: 60089
                                                                                                                                                                                                                                                    COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                         ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: CUCUMBER MOSAIC VIRUS
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3..660
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Matches:
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Indels:
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                                                                                                                                         Version #1.30
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Score: Pred. No.:

RESULT 4

ATTORNEY/AGENT INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5400

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UŞ-09-857-841-4 (1-14) x US-10-437-963-3120 (1-792)
                                                   Percent Similarity:
Best Local Similarity:
Query Match:
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Best Local Similarity:
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                                                                                                                                                                                                                   APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Wu, Wei
APPLICANT: Wu, Wei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Boukharov, Andrey A.
APPLICANT: Li, Ping
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 3120
LENGTH: 792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 3120, Application US/10437963
Publication No. US20040123343A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
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                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Oryza sativa
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 960 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORIGINAL SOURCE:
ORGANISM: CUCUMBER MOSAIC VIRUS
STRAIN: STRAIN C
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LOCATION: 1..658
SEQUENCE DESCRIPTION:
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US-10-424-599-140904
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Best Local Similarity:
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  Percent Similarity:
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                                                                                                                                                                                                                                                                             APPLICANT: LA ROSA, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(5322)B
FILE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 66023
LENGTH: 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 140904
LENGTH: 107
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APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)B
FULL REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
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                                                                                                                FEATURE: FEATURE: Clone ID: MRT4577_160212C.1
                                                                                                                                                            FEATURE:
NAME/KEY: unsure
LOCATION: (1)...(166)
OTHER INFORMATION: unsure at all n locations
                                                                                                                                                                                                                                            ORGANISM: Zea mays
                                                                                                                                                                                                                                                                   TYPE: DNA
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ORGANISM: Glycine max
FEATURE:
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Matches:
Conservative:
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US-10-424-599-94618
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Query Match:
                                          Pred. No.:
                                                           Alignment Scores:
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DB:
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    Percent Similarity:
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CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 94618
LENGTH: 233
                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 15112
SEQ ID NO 7184
                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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Patent No. US20020137139A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 94618, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION:
                                                                                                                                                                                                                              APPLICANT: Warren, Wesley C.
APPLICANT: Tao, Nengbing
APPLICANT: Tao, Nengbing
APPLICANT: Byatt, John C.
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILL REFERENCE: 16511.006/37-21 (10298) C
CURRENT APPLICATION NUMBER: US/09/960,352
CURRENT FILING DATE: 2001-09-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: La Roba Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
                                                                                                    TYPE: DNA
ORGANISM: Bos taurus
OTHER INFORMATION: Clone ID: 31-LIB3057-014-Q1-K1-H3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Glycine max
                                                                                                                                                                             LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         No . .
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                                                                                                                                                                                 279
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Best Local Similarity:
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US-10-425-115-46484
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                                                                             Alignment Scores:
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 34460
LENGTH: 408
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SEQ ID NO 46484
LENGTH: 316
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APPLICANT: La Rosa, Thomas
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                                                                                                                                                                                                                                               APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-756
CURRENT APPLICATION NUMBER: US/09/918,995
CURRENT FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: US/09/235,076
PRIOR FILING DATE: 1999-01-20
PRIOR FILING DATE: 1999-01-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
                                                                                                                                    TYPE: DNA ORGANISM: Homo sapiens
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ORGANISM: Zea mays
FEATURE:
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; ORGANISM: Homo sapiens
US-09-998-598-1555
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US-09-998-598-1555/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15219, Application US/09864761 Patent No. US20020048763A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 1555
LENGTH: 452
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                                                                                                  TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TIVELE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY

FILE REFERENCE: Acomica-X-1

CURRENT APPLICATION NUMBER: US/09/864,761

CURRENT FILING DATE: 2001-05-23

PRIOR APPLICATION NUMBER: US 60/180,312

PRIOR APPLICATION NUMBER: US 60/180,312

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR APPLICATION NUMBER: US 60/32,366

PRIOR FILING DATE: 2000-08-03

PRIOR APPLICATION NUMBER: US 60/32,366

PRIOR APPLICATION NUMBER: US 60/32,366

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR PILING DATE: 2000-09-27

PRIOR PILING DATE: 2000-09-27
PRIOR
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APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
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APPLICANT: Xu, Jiangchun
APPLICANT: Chenault, Ruth A.
APPLICANT: Chenault, Ruth A.
APPLICANT: Meagher, Madelein Joy
APPLICANT: MEAGHER, MACHEIN JOY
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSTIONS AND METHODS
FILE REFERENCE: 210121.561
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SOFTWARE: Corixa Invention Disclosure Database
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CURRENT FILING DATE: 2001-11-16
                          APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
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Best Local Similarity:
Query Match:
                                                                       ORGANISM: Zea mays
FEATURE:
OTHER INFORMATION: Clone ID: MRT4577_90708C.1
US-10-425-115-174482
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US-10-425-115-174482/c
                   Pred. No.:
                                  Alignment Scores:
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; OTHER INFORMATION: EXPRESSED IN FETAL LIVER,
US-09-864-761-15219
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CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 174482
LENGTH: 478
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SEQ ID NO 15219
LENGTH: 464
                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                           APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 49117
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ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                  TYPE: DNA
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FILING DATE: 2000-06-30
APPLICATION NUMBER: US 09/774,203
FILING DATE: 2001-01-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
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APPLICATION NUMBER: PCT/US01/00661
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FILING DATE: 2001-01-30
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1: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
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Copyright (c) 1993 - 2004 Compugen Ltd.
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US-10-025-915-1131
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272	263	251	250	240	223	202	199	176	171	168	160	156	142	135	131	127	126	107	105	100	98	91	85	83	83	81	78	83	65	54	15
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-10-425-115-2870	-10-408-765A-260	-10-425-114-3911	-10-425-115-2356	-10-282-122A-7045	-10-369-493-1984	-10-437-963-11073	-10-437-963-1961	-10-425-115-28451	-10-425-115-33096	-10-425-114-3943	-10-425-115-26107	-437-963-121	-10-437-963-11073	-10-425-114-5830	-10-437-963-10565	-10-437-963-1248	-10-425-114-5907	-10-425-115-29123	-10-437-963-17762	0-437-963-1032	-10-425-115-35695	-10-425-115-24405	-10-425-115-2056	-10-425-115-31123	-10-424-599-14532	-10-425-115-35282	-10-424-599-25638	-10-425-115-28841	25-115-3111	-10-424-599-14541	15A-1
nce 287096,	equence 2608,	0	equence 235635,	equence 70451,	e	equence 11073	equence 19617	e 28451	equence :	e 39432	equence 26107	Sequence 121791,	11073	58305	e 10565		e 59079	equence 29123	equence 17762	e _	equence 35695	e 24405	e 20560	nce 31123	nce 14532	nce 35282	equence 25638	nce 28841	e 31117	equence 14541	Sequence 133, App

## ALIGNMENTS

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US-10-011-033-2

US-10-011-033-2

Sequence 2, Application US/10011033

Publication No. US20020124286A1

GENERAL INFORMATION:

APPLICANT: Boeshore, Maury L

MCMaster, J. Russell

Tricoli, David M

Reynolds, John F

Carney, Kim J

COLUMBER OF INVENTION: Plants Resistant to C Strains of

CUcumber Mosaic Virus

NUMBER OF SEQUENCES: 15

CORRESPONDENCE ADDRESS:

ADDRESSEE: Rockey, Milnamow & Katz, Ltd.

STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700

CITY: Chicago

STATE: Illinois

COUNTRY: U. S. A.

ZIP: 60089

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER PAPLICATION NUMBER: US/08/MS-DOS

SOFTMARE: Datentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/011,033

FILING DATE: 13-No. US20020124286A1-2001

CLASSIFICATION NUMBER: US/08/875,233

FILING DATE: 26-UN-1997

ATTORNEY/AGENT INFORMATION:

TELEPHONE: 312-616-5400
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LENGTH: 218 amino acids
TYPE: amino acid
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
US-10-011-033-4
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                                        Best Loc
Matches
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                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/10011033 Publication No. US20020124286A1
                                                                                                                                                                                                                               TELEFAX: 312-616-5460 INFORMATION FOR SEQ ID NO: 4: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/011,033

FILING DATE: 13-No. US20020124286A1-2001

CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/875,233

FILING DATE: 26-JUN-1997

ATTORNEY/AGENT INFORMATION:

NAME: Lisa V. Mueller

TELECOMMUNICATION INFORMATION:

TELEPHONE: 1312-616-6400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 60089

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS: ADDRESSEE: ROckey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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CITY: Chicago
STATE: Illinois
MDKSESTSAGRNRR 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 218 amino acids
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                                        100.0%; Score 14; DB 13; larity 100.0%; Pred. No. 2.2e-06; Conservative 0; Mismatches 0;
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: 15
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Pred. No. 2.2e-06;
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                                                                             Length 218;
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; MOLECULE TYPE: protein ; SEQUENCE DESCRIPTION: SEQ ID NO: 6: US-10-011-033-6
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                                                                                                                                                                                                                                                   Sequence 15, Applicati
Publication No. US2002
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 6, Application US/10011033
Publication No. US20020124286A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 312-616-5460 INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/011,033
APPLICATION NUMBER: US/0020124286A1-2001
ETILING DATE: 13-NO. US20020124286A1-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-54000
                                                                                APPLICANT: Boeshore, Maury L
McMaster, J. Russell
Tricoli, David M
Tricoli, John F
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C
Cucumber Mosaic Virus
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Tricoli, David M
Reynolds, John F
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
CORRESSES: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Boeshore, Maury L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ch 100.0%;
l Similarity 100.0%;
l4; Conservative 0;
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
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STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson,
                                                                                                                                                                                                                                                                         Application US/10011033 No. US20020124286A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 218 amino acids
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Pred. No. 2.2e-06;
); Mismatches 0;
                                                                                                               Strains
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/10/011,033
FILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION ATA:
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
TELEPHONE: 312-616-5460
TELEPAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acids
LENGTH: 218 amino acids
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US-10-011-033-10
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Publication No. US2002
GENERAL INFORMATION:
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Best Local Similarity
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TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTMARE: PAtentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                   APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lisa V. Mueller
                                                                                             APPLICATION NUMBER: US/10/011,033
PILING DATE: 13-No. US20020124286A1-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Reynolds, John F
Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains of
                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Rockey, Milnamow & Katz, Ltd.
STREET: 2 Prudential Plaza, 180 N. Stetson, Suite 4700
CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A.
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McMaster, J. Russell
Tricoli, David M
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                John F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
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Query Match
Best Local Similarity
7; Conserve
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US-10-424-599-283746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-425-114-51049
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: LA ROSA Thomas J
APPLICANT: Kovalic David K
APPLICANT: Chou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)B
FULE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 283746
LENGTH: 35
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                                                  APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Xovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 51049
LENGTH: 183
TWORLD TO THE TABLE TABLE TO THE TABLE 
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Publication No. US20040034888A1
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Publication No. US20040031072A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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ORGANISM: Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
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TYPE: amino acid
TOPOLOGY: linear
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100.0%; Pred. No. 4.9;
tive 0; Mismatches
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APPLICANT: Kovalic, David K.

APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 53499
LENGTH: 343
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                    RESULT 9
WS-10-425-115-323368
; Sequence 323368, Application US/10425115
; Publication No. US20040214272A1
Query Match
Best Local Similarity
7; Conserve
                                                                            ; OTHER INFORMATION: Clone ID: MRT4577_57983C.1.pep US-10-425-115-323368
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; OTHER INFORMATION: Clone ID: 700836661_FLI.pep
US-10-425-114-53499
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SEQ ID NO 323368
LENGTH: 375
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(3322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
                                                                                                            FEATURE:
                                                                                                                           ORGANISM: Zea mays
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             / Match 50.0%; Score 7; DB:
Local Similarity 100.0%; Pred. No. 19
1es 7; Conservative 0; Mismatches
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                                             50.0%;
           Score 7; DB 17; Length 375; Pred. No. 34; 0; Mismatches 0; Indels
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US-10-424-599-236497; Sequence 236497, Application US/10424599; Publication No. US20040031072A1
                                                                                                ; OTHER INFORMATION: Clone ID: PAT_MRT3847_55584C.1.pep
US-10-424-599-236497
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US-10-437-963-169865
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US-10-437-963-169865
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APPLICANT: Kovalic David K
APPLICANT: Can Yongwei
TITILE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITILE OF INVENTION: Blants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
FULE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 236497
LENGTH: 469
TUREN DET
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Query Match 50.0%; Score 7; DB 15; Length 469; Best Local Similarity 100.0%; Pred. No. 41; Matches 7; Conservative 0; Mismatches 0; Indels
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SEQ ID NO 169865
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CURRENT FILING DATE: 2003-05-14
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APPLICANT:
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                                                                                                                                             ORGANISM: Glycine max FEATURE:
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OTHER INFORMATION: unsure at all Xaa locations
FEATURE:
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Zhou, Yihua
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                                                                              RESULT 14
US-10-203-915A-133
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TITLE OF INVENTION: Method for Antigen-Specific Stimulation of T
TITLE OF INVENTION: Lymphocytes with synthetic Peptide Libraries
FILE REPERENCE: 010266wc/JH/ml
CURRENT APPLICATION NUMBER: US/10/203,915A
CURRENT FILING DATE: 2002-08-15
NUMBER OF SEQ ID NOS: 260
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 131
                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Method for Antigen-Specific Stimulation of T
TITLE OF INVENTION: Lymphocytes with synthetic Peptide Libraries
FILE REFERENCE: 010266wo/JH/ml
CURRENT APPLICATION NUMBER: US/10/203,915A
CURRENT FILING DATE: 2002-08-15
NUMBER OF SEQ ID NOS: 260
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 132
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE: NECESSAUTION. Description of Artificial Sequence
FEATURE: NECESSAUTION.
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                Sequence 133, Application US/10203915A Publication No. US20040106159A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 132, Application No. US200 GENERAL INFORMATION: APPLICANT: Kern, F1
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Best Local 9
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APPLICANT: Kern,
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nes 6; Conserv
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Local Similarity 100.0%; Pred. No.
1es 6; Conservative 0; Mismatch
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o. US20040106159A1
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100.0%; Pr
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100.0%; Pred. No. 24
Live 0; Mismatches
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o. 24;
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Search completed: December Job time : 67 secs

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; OTHER INFORMATION: Description of Artificial Sequence: HCMV pp65; OTHER INFORMATION: Fragment US-10-203-915A-133
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                                                                                                                                                                                                                                          Sequence 14541, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
APPLICANT: LA ROSA Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITLE OF INVENTION UNMEER: US/10/424,599
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 145411
LENGTH: 54
                                                                                                                                                       ; OTHER INFORMATION: Clone ID: PAT_MRT3847_102325C.1.pep
US-10-424-599-145411
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Matches
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TITLE OF INVENTION: Lymphocytes with synthetic Peptide Libraries
FILE REFERENCE: 010266wo/JH/ml
CURRENT APPLICATION NUMBER: US/10/203,915A
CURRENT FILING DATE: 2002-08-15
NUMBER OF SEQ ID NOS: 260
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 133
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Best Local
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Best Local
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## ALIGNMENTS

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Of6133 PRELIMINARY; PRT; 217 AA.  O66133, O1-NOV-1996 (TrEMBLrel. 01, Created) O1-NOV-1996 (TrEMBLrel. 01, Last sequence update) O1-OCT-2003 (TrEMBLrel. 25, Last annotation update) Coat protein. Cucumber mosaic cucumovirus). Viruses; ssRNA positive-strand viruses, no DNA stage; Bromoviridae; Cucumovirus. NCBL_TaxID=12305; [1] SEQUENCE FROM N.A. Reichel H., Marino L., Kummert J., Belalcazar S., Narvaez J.; "Caracterizacion del gen de la proteina de la capside de dos aislamientos del virus del mosaico del pepino (CMV), obtenidos de platano y banano (Musa spp.).";	Query Match 100.0%; Score 14; DB 2; Length 217; Best Local Similarity 100.0%; Pred. No. 3.6e-07; Matches 14; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  1 MDKSESTSAGRNRR 14	OGTTD4 PRELIMINARY; PRT; 217 AA.  OGTTD4,  OS-JUL-2004 (TrEMBLrel. 27, Created)  OS-JUL-2004 (TrEMBLrel. 27, Last sequence update)  OS-JUL-2004 (TrEMBLrel. 27, Last annotation update)  OS-JUL-2004 (TrEMBLrel. 27, Last annotation update)  Cout protein.  Cucumber mosaic virus (cucumber mosaic cucumovirus).  Viruses; seRNA positive-strand viruses, no DNA stage; Bromoviridae;  Cucumovirus.  NCBI_TAXID=12305;  [1]  SEQUENCE FROM N.A.  Yan L., Xu Z., Goldbach R., Chen K., Prins M.;  SEQUENCE FROM S.,  Submitted (CCT-2003) to the EMBL/GenBank/DDBJ databases.  EMBL; AY429437; AAR89478.1; -  GO; GO:0005198; F:structural molecule activity; IEA.  GO; GO:0005198; Cucumo_coat; 1.  PRINTS; PR00760; Cucumo_coat; 1.  PRINTS; PR00722; CUCUMOCOAT.  PRODOM; PD001284; Cucumovirus_coat; 1.  Coat protein.  SEQUENCE 217 AA; 24015 MM; 32D0035B4F20E891 CRC64;

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InterPro; IPR000247; Cucumovirus coat.

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GO; GO:0005198; F:structural molecule activity;
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EMBL; U32859; AAB50176.1;
PIR; JC6073; JC6073.
PIR; JC6075; JC6075.
                                                                                                                                                                                AAR89478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Daolin D., Jie S., Peng Z., Zhixin L., Xiaodong D., "Cloning and Reconstruction of a Expression Vector i Mosaic Virus Coat Protein Gene.";
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Banana mosaic virus.
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                               Cucumber mosaic virus (cucumber mosaic cucumovirus)
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Pred. No. 3.6e-07;
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Hyon S., Park Y.I.;
Submitted (FEB-1994) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: Belongs to the cucumoviruses/bromoviruses coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Viruses; ssRNA positive-strand viruses, 1 Cucumovirus.
                                                                                                                                                                      PIR; $42098; $42098; Cucumovirus coat.
InterPro; IPR000247; Cucumovirus coat.
InterPro; IPR0008975; Viral_cap_coat.
Pfam; PF00750; Cucumo coat; 1.
PRINTS; PR00222; CUCUMOCOAT.
ProDom; PD001284; Cucumovirus coat; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Coat protein (Capsid protein)
Cucumber mosaic virus (strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (OCT-2003) to the EMBL/GenBank/DDBJ databases EMBL; AY429437; AAR89478.1; -. Coat protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yan L., Xu Z., Goldbach R., "Complete nucleotide sequenc RNA3.";
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                                                                                                                                          Coat protein.
SEQUENCE 21
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(Rel. 37, Last annotation update)
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                      entities
or send a
                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
Chaumpluk P., Sasaki Y., Nakajima N., Nagano H., Nakamura I.,
Suzuki K., Mise K., Inouye N., Okuno T., Furusawa I.;
Sizuki K. subgroup I members of Japanese cucumber mosaic virus
determined by nucleotide sequence analysis on RNA3's cDNAs.";
Ann. Phytopathol. Soc. Jpn. 62:40-44(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation updat
Coat protein (Capsid protein) (CP).
Cucumber mosaic virus (strain C7-2) (CMV).
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"Natural infection of banana by a satellite-containing strain of
cucumber mosaic virus.";
Phytoparasitica 24:49-56(1996).
-!- SIWILARITY: Belongs to the cucumoviruses/bromoviruses coat protein
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                        This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Cucumber mosaic virus (strain Banana) (CMV).
Viruses; ssRNA positive-strand viruses, no DNA stage; Bromoviridae;
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InterPro; IPR008975; Viral_cap_coat.
Pfam; PF00760; Cucumo_coat; 1.
PRINTS; PR00222; CUCUMOCOAT.
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NCBI_TaxID=117120;
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SEQUENCE 218 AA;
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ProDom; PD001284; Cucumovirus_coat; 1.
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RESULT 9
COAT\_CMVFC

54555

01-DEC-1992 01-DEC-1992 15-DEC-1998

(Rel. 24, Created) (Rel. 24, Last seq (Rel. 37, Last ann

sequence up

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COAT CMVFC

STANDARD;

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                                                                   Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q66143;
15-DEC-1998
15-DEC-1998
15-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                      Chaumpluk P., Sasaki Y., Nakajima N., Nagano H., Nakamura I., Suzuki K., Mise K., Inouye N., Okuno T., Furusawa I., "Six new subgroup I members of Japanese cucumber mosaic virus determined by nucleotide sequence analysis on kNA3's CDNAs.", Ann. Phytopathol. Soc. Jpn. 62:40-44(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CMVCS
                                                                                                                                                   EMBL; D28489; BAA05851.1; -.
InterPro; IPR000247; Cucumovirus coat.
InterPro; IPR008975; Viral cap coat.
Pfam; PF00760; Cucumo_coat; 1.
PRINTS; PR00222; CUCUMOCOAT.
Dropos BR001284; CUCUMOCOAT.
                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMBL outst the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for commodified and this statement is not removed. Usage by and for commodified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Coat protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; D42079; BAA07675.1; -.
InterPro; IPR000247; Cucumovirus coat.
InterPro; IPR008975; Viral_cap_coat.
Pfam; PF00760; Cucumo coat; 1.
PRINTS; PR00222; CUCUMOCOAT.
ProDom; PD001284; Cucumovirus_coat; 1.
                                                                                                               Coat protein
SEQUENCE 2:
                                                                                                                                                                                                                                            entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cucumovirus.
NCBI_TaxID=117109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Coat protein (Capsid protein) (CP Cucumber mosaic virus (strain CS)
                                                                                                                                       ProDom; PD001284; Cucumovirus_coat; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Viruses; ssRNA positive-strand viruses,
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                                                                                                                                                                                                                                                                                                                                                                             SIMILARITY:
                                                                                                                                                                                                                                                                                                                                                                 family.
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                                                      14;
                                                                     Similarity
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                  MDKSESTSAGRNRR 14
MDKSESTSAGRNRR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MDKSESTSAGRNRR 14
                                                                                                               218 AA;
                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (Rel. 37, Created)
(Rel. 37, Last sequence update)
(Rel. 37, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STANDARD;
                                                                                                                                                                                                                                                                                                                                                                             Belongs
                                                                                                                                                                                                                                                                                                                                                                       Soc. Jpn. 62:40-44(1996).
elongs to the cucumoviruses/bromoviruses coat protein
                                                                                                               24157 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24033 MW;
                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 14; Di
100.0%; Pred. No. 3.
tive 0; Mismatches
                                                      <u>.</u>
                                                      Score 14; DE
Pred. No. 3.7
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CP)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16D96022D1A26FD8 CRC64;
                                                                                                               76F6BDCD9628F683 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CMV).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                218
                                                                                                                                                                                                                                                             moved. Usage by and fo
(See http://www.isb-sib.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           no DNA stage; Bromoviridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.7e-07;
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                                                                                    DB 1;
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                                                                     .7e-07;
                                                                                   Length 218;
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                                                                                                                                                                                                                                                                                                                                    a collaboration
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Best Local S
Matches 14
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Q66140;
15-DEC-1998
                                                                                              EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.

Chaumpluk P., Sasaki Y., Nakajima N., Nagano H., Nakamura I.,

Chaumpluk P., Sasaki Y., Inouye N., Okuno T., Furusawa I.;

Suzuki K., Mise K., Inouye N., Okuno T., Furusawa I.;

"Six new subgroup I members of Japanese cucumber mosaic virus as

determined by nucleotide sequence analysis on RNA3 s cDNAs.";

determined by nucleotide sequence analysis on RNA3 s cDNAs.";

Ann. Phytopathol. Soc. Jpn. 62:40-44(1996).

-I- SIMILARITY: Belongs to the cucumoviruses/bromoviruses coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Coat protein (Capsid protein) (CP).
Cucumber mosaic virus (strain FT) (CMV).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PIR; JQ1253; JQ1253. Cucumovirus coat. InterPro; IPR000247; Cucumovirus coat. InterPro; IPR008975; Viral_cap_coat. Pfam; PF00750; Cucumo coat; 1. PRINTS; PR00222; CUCUMOCOAT. PRODom; PD001284; Cucumovirus_coat; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                               muse by non-profit institutions as long a modified and this statement is not removed. entitles requires a license agreement (See lor send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its the by non-profit institutions as long as its content is in no way
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MEDLINE=92013983; PubMed=1919534;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Viruses; ssRNA positive-strand viruses, Cucumovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Viruses; seRNA positive-strand viruses, no DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shintaku M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Coat protein (Capsid protein)
Cucumber mosaic virus (strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                induction
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                                                                                                      D28487;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      D10544; BAA01403.1; -.
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                                                                                                  BAA05847.1;
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virus (strain FC) (CMV).
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                                                       Cucumovirus_coat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              change correlating with chlorosis
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
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                                                                                                                                                                                                                          (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.7e-07;
hes 0;
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                                                                                                                                                                                                                                                                 Usage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           stage; Bromoviridae;
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RESULT
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Query Match
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                                                              EMBL; X16386; CAA34422.1; -.
EMBL; D10538; BAA01397.1; -.
PIR; S09663; S09663; CUCUMOVÍTUS COAT.
InterPro; IPR000247; CucumovíTus Coat.
Pfam; PF00760; Cucumo Coat; 1.
PRINTS; PR00222; CUCUMOCOAT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P14767;
01-APR-1990
01-APR-1990
15-DEC-1998
                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Ew the European Bioinformatics Institute. There are no restrues by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entitles requires a license agreement (See http://www.isb-s or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=90098885; PubMed=2602158;
Noel M.J.T., Ben Tahar S.;
"Nucleotide sequence of the coat protein gene cucumber mosaic virus (CMV) strain I17F.";
Nucleic Acids Res. 17:10492-10492(1989).
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01-APR-1990 (Rel. 14, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Coat protein (Capsid protein) (CP).
Cucumber mosaic virus (strain I17F) (CMV), an
Cucumber mosaic virus (strain FNY) (CMV).
                                      ProDom; PD001284; Cucumovirus_coat;
Coat protein.
                                                                                                                                                                                                                                                                                                         J. Gen.
                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=90206832; PubMed=2320436;
Noel M.J.T., Ben Tahar S.;
Nucleic Acids Res. 18:1332-1332(1
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PRINTS; PR00222; CUCUMOCOAT.
ProDom; PD001284; Cucumovirus_coat;
                                                                                                                                                                                                                                                                                                                    "Nucleotide sequence and evolutionary relationships virus (CMV) strains: CMV RNA 3.";
                                                                                                                                                                                                                                                                                                                                               MEDLINE=91037954; PubMed=2230731;
Owen J., Shintaku M., Aeschleman P.,
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Pfam; PF00760; Cucumo_coat; 1.
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SIMILARITY: Belongs to the cucumoviruses/bromoviruses coat protein
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s. 18:1332-1332(1990).
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Score 14;
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15-DEC-1998 (Rel. 37, I
15-DEC-1998 (Rel. 37, I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Karasawa A., Ito A., Okada I., Hase S., Ehara Y.;
"A possible role of RNA 2 of cucumber mosaic cucumovirus as a
determinant of infection phenotype on cowpea.";
Ann. Phytopathol. Soc. Jpn. 63:289-297(1997).
-!- SIMILARITY: Belongs to the cucumoviruses/bromoviruses coat protein
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15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update
15-DEC-1998 (Rel. 37, Last annotation update
Coat protein (Capsid protein) (CP).
Cucumber mosaic virus (strain lizuka) (CMV).
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Paek K.H.;
Submitted (SEP-1994) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: Belongs to the cucumoviruses/bromoviruses coat protein
                                                                                                                                                                                               15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Coat protein (Capsid protein) (CP).
Cucumber mosaic virus (strain Kor) (CMV).
                                                SEQUENCE FROM N.A.
Kim S.J., Cho H.S.,
Paek K.H.;
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InterPro; IPR008975; Viral_cap_coat.
Pfam; PF00760; Cucumo_coat; 1.
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SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
MEDLINE=91037954; PubMed=2230731;
MEDLINE=91037954; PubMed=2230731;
Owen J., Shintaku M., Aeschleman P., Tah;
"Nucleotide sequence and evolutionary re
r virus (CMV) strains: CMV RNA 3.";
T Gen. Virol. 71:2243-2249(1990).
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01-DEC-1992
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                                                                                                                                                                          InterPro; IPR000247; Cucumovirus_coa
InterPro; IPR008975; Viral_coat.
Pfam; PF00760; Cucumo_coat; 1.
PRINTS; PR00222; CUCUMOCOAT.
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-!- SIMILARITY: Belongs to the cucumoviruses/bromoviruses coat
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Pfam; PF00750; Cucumo coat; 1.
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                                                                                                                          Coat protein.
SEQUENCE 218
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MDKSESTSAGRNRR 14

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COAT_CMVN
ID COAT_CMVN
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ID COAT_CMVN
AC Q66138;
DT 15-DEC-1998 (Rel. 37, Last sequence update)
DT 15-DEC-1998 (Rel. 37, Last annotation update)
DE Coat protein (Capsid protein) (CP).
CO Viruses; ssRNA positive-strand viruses, no DNA st
Coutmovirus.

OX NCBI_TAXID=117123;
RN [1]
RP SEQUENCE FROM N.A.
RA SUZUKİ K., Mise K., Inoue N., Okuno T., Furusaw,
RT SEQUENCE FROM N.A.
SAZUKİ K., Mise K., Inoue N., Okuno T., Furusaw,
RT determined by nucleotide sequence analysis on RNJ
RA SUZUKİ K., Mise K., Inoue N., Okuno T., Furusaw,
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RA SUZUKİ K., Mise K., Inoue N., Okuno T., Furusaw,
RT determined by nucleotide sequence analysis on RNJ
RA SUZUKİ K., Mise K., Inoue N., Okuno T., Furusaw,
RT SIMILARITY: Belongs to the cucumoviruses/brot
CC family.

CC -1-SIMILARITY: Belongs to the cucumoviruse.
CC family.

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InterPro; IPR000247; Cucumovirus coat.
InterPro; IPR000975; Viral_cap_coat.
Pfam; pF00760; Cucumo_coat; 1.
PRINTS; PR00222; CUCUMOCOAT.
PRINTS; PR001284; Cucumovirus_coat; 1.
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SEQUENCE 218 AA; 24078 MW; 0BA67C84D8F41B22 CRC64;
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Chaumpluk P., Sasaki Y., Nakajima N., Nagano H., Nakamura I.,
Chaumpluk P., Sasaki Y., Nakajima N., Nagano H., Nakamura I.,
Chaumpluk P., Sasaki Y., Nakajima N., Nagano H., Nakamura I.,
Suzuki K., Mise K., Inouye N., Okuno T., Furusawa I.;
Suzuki K., Mise K., Inouye N., Okuno T., Purusawa I.;
"Six new subgroup I members of Japanese cucumber mosaic virus as
determined by nucleotide sequence analysis on RNA3's cDNAs.";
Ann. Phytopathol. Soc. Jpn. 62:40-44(1996).

Ann. Phytopathol. Soc. Jpn. 62:40-44(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Viruses; ssRNA positive-strand viruses, no DNA stage; Bromoviridae;
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US-08-482-663-36	US-08-090-192-36	US-08-488-027-36	US-08-486-569-36	US-08-488-031-36	US-08-090-193-36	US-08-615-181-93	US-09-134-001C-4463	US-09-252-991A-30085	US-09-252-991A-32419	US-09-252-991A-18657	US-09-194-468A-30	US-09-248-796A-19191	US-08-605-541B-12	US-09-171-699-8	US-09-171-699-6	US-09-252-991A-23246	US-09-812-079A-2
Sequence 36	Sequence 36	Sequence 36	Sequence 36	Sequence 36	Sequence 36	Sequence 93,	Sequence 44	Sequence 30	Sequence 32	Sequence 18	Sequence 30	Sequence 19	Sequence 12	Sequence 8,	Sequence 6,	Sequence 23	Sequence 2,
6, Appl	6, Appl	•	6, Appl	6, Appl	-	3, Appl	4463, Ap	30085, A	32419, A	18657, A	30, Appl	9191, A	2, Appl	, Appli	•	23246, A	, Appli

## ALIGNMENTS

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RESULT 1
542259-2
;PACENT NO. 542259
;PACENT NO. 5422259
;PACENT NO. 5422259
; PAPLICANT: DE BOTH, MICHIEL;BEN TAHAR, SOPHIA;NOEL, MARIANNE;
;PERRET, JOEL
; CITILE OF INVENTION: TRANSGENIC PLANTS BELONGING TO THE
;SPECIES CUCUMIS MELO
; NUMBER OF SEQUENCES: 4
; CURRENT APPLICATION NUMBER: US/08/27,563
; PILING DATE: 05-MAR-1993
; PRICR APPLICATION NUMBER: 566,255
; APPLICATION NUMBER: 566,255
; PILING DATE: 13-AUG-1990
; PILING DATE: 13-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-398-209-2
                                                                                                                                                                                                                                                                                                                                                       Patent No. 5789656
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/08398209 Patent No. 5789656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                     APPLICANT: Ben Tahar, Sophia
APPLICANT: Perret, Joel
TITLE OF INVENTION: Transgenic Plants Belonging
TITLE OF INVENTION: Species Cucumis Melo
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Merchant & Gould
STREET: 90 South 7th Street, 3100 No. 5789656w
COUNTRY: USA
ZIP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                APPLICANT: DeBoth, Michiel APPLICANT: No. 57896561, Marianne
                                                                                                                              STREET: 90 South
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14;
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Pred. No. 2.6e-07;
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                                                                                                                                                           5789656west Center
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                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: DeBoth, Michiel
APPLICANT: No. 57896561, Marianne
APPLICANT: Ben Tahar, Sophia
APPLICANT: Perret, Joel
TITLE OF INVENTION: Transgenic Plants Belonging to the
TITLE OF INVENTION: Species Cucumis Melo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 2:
            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/027,563
PILING DATE: 05-MAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/566,255
                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NATA:
APPLICATION NUMBER: US 08/027,563
APPLICATION NUMBER: US 08/027,563
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/566,255
FILING DATE: 13-AUG-1990
                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSE: Merchant & Gould
STREET: 90 South 7th Street, 3100 No. 5789656west Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 11-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 218 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Hillson, Randall A.
REGISTRATION NUMBER: 31,838
REFERENCE/DOCKET NUMBER: 80
APPLICATION NUMBER: US 07 FILING DATE: 13-AUG-1990
                                                                                                                                      FILING DATE:
                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                    STREET: 90 South CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION FILING DATE:
                                                                                                                                                                                                                                                                                                                   COUNTRY:
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                                                                                                                                                                                                                                                                                                 55402
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                                                                                                                                                                                                                                                                                                                   USA
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US-08-553-619B-3
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                                   Matches
                                                  Query Match
Best Local &
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                                                                                                                                                                                                                                    CLASS FICATION: 800
CLASS FICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: MATCUS-WYDEY, LYRN
REGISTRATION NUMBER: 34,869
REFERENCE/DOCKET NUMBER: 137-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/354-3588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: DeHaan
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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                  TELEFAX: 415/857-1125
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: DeHaan, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: protein
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/553,619B FILING DATE: December 1, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U
ZIP: 94304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 975 Call
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TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 31,838
REFERENCE/DOCKET NUMBER: 8076.21US02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: No. 5919705artis Crop Protection STREET: 975 California Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14;
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                                   14;
 1 MDKSESTSAGRNRR 14
                                                    Similarity
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                                                                                                                                                                      218 amino acids
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                                   Conservative
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Virus Resistant Plants
                                                  100.0%; Score 14; DB 2; 100.0%; Pred. No. 2.6e-07;
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Pred. No. 2.6e-07;
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                                    Mismatches
                                                                Length 218;
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                                   Indels
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                                   Gaps
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MDKSESTSAGRNRR 14

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; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-875-233-2
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US-08-875-233-2
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                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/08875233
Patent No. 6127601
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.0%; Score 14; DB 3; Best Local Similarity 100.0%; Pred. No. 2.6e-07, Matches 14; Conservative 0; Mismatches 0.
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Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: FORM:
MEDIUM TYPE: FORM:
COMPUTER: IBM PC compatible
COMPUTER: PATEN: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/08/875,233
FILING DATE: 26-UN-1997
CLASSIFICATION: 800
CCCNESSIFICATION: 800
                                                                                                                                                                                APPLICANT: Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
TITLE OF INVENTION: Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                          APPLICANT: Boeshore, Maury L
APPLICANT: McMaster, J. Russell
APPLICANT: Tricoli, David M
APPLICANT: Reynolds, John F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: 312-616-5400
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OMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Lisa V. Mueller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TLE OF INVENTION: Plants Resistant to C Strains
                                                            STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Chicago
STATE: Illinois
                                                                                                                    ADDRESSEE:
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                                  COUNTRY:
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                                                       Chicago
Illinois
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    Rockey, Milnamow & Katz, Ltd.
    Prudential Plaza, 180 N. Stetson, Suite 4700

                                                                                                3: Rockey, Milnamow & Katz, Ltd.
2 Prudential Plaza, 180 N. Stetson, Suite 4700
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Tricoli, David M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Boeshore, Maury L
                                    U.S.A.
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US-08-875-233-6
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TELECOMMUNICATION INFORMATION:
TELEPHONB: 312-616-5400
TELEFAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local Similarity
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                                                                   TELEPAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 218 amino acids
                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC COMPATION
SOPTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/875,233
FILING DATE: 26-UN-1997
CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/875,233
FILING DATE: 26-UN-1997
CLASSIFICATION: 800
CLASSIFICATION: 800
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APPLICANT:
APPLICANT:
                                                                                                                                                                NAME: Lisa V. Mueller TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Chicago
STATE: Illinois
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE:
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                                    TOPOLOGY:
                                                                                                                                             TELEPHONE:
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                                                  amino acids
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2 Prudential Plaza, 180 N. Stetson, Suite 4700
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McMaster, J. Russell
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                                                                                                                                               312-616-5400
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100.0%; Pred. No. 2.6e-07;
rative 0; Mismatches 0;
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Query Match

100.0%;

Score 14;

DB 3;

Length 218;

Best Local Similarity

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RESULT 8
US-08-875-233-15
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                                                                      GENERAL INFORMATION:
APPLICANT: DE BOTH, MICHAEL
APPLICANT: NOEL, MARIANNE
APPLICANT: TAHAR, SOPHIA BEN
APPLICANT: PERRET, JOEL
APPLICANT: BIOSEM
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US-09-127-742-2
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                   Matches
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TITLE OF INVENTION: TRANSGENIC PLANTS BELONGING TO THE SPECIES CUCUMIS MELO FILE REFERENCE: 8076.21USD.2 CURRENT APPLICATION INVMER: US/09/127,742A CURRENT FILING DATE: 1998-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
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APPLICANT:
APPLICANT:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION WIMBER: US/08/875,233
FILING DATE: 26-UN-1997
CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: Chicago
CITY: Chicago
TTATE: Illinois
TT.S.A.
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NAME: Lisa V. Mueller
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Bosehore, Maury L
APPLICANT: McMaster, J. Russell
APPLICANT: Tricoli, David M
APPLICANT: Reynolds, John F
APPLICANT: Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
TITLE OF INVENTION: Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U.S.A. ZIP: 60089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 218 amino acids
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    Rockey, Milnamow & Katz, Ltd.
    Prudential Plaza, 180 N. Stetson, Suite 4700

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RESULT 11
5422259-4
; Patent No. 5422259
; APPLICANT: DE BOTH, MICHIEL; BEN TAHAR, SOPHIA; NOEL, MARIANNE;
; ERRET, JOEL
; PERRET, JOEL
; TITLE OF INVENTION: TRANSGENIC PLANTS BELONGING TO THE
; SPECIES CUCCUMIS MELO
; PUMBER OF SEQUENCES: 4
; CURRENT APPLICATION DATA:
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NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
LENGTH: 218
TYPE: PRT
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US-09-127-742-4
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SEQ ID NO 4
LENGTH: 218
TYPE: PRT
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4, Application US/09127742A Patent No. 6198022
                                                                                                                                                                                                                                                                                                                                                                                         Matches
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APPLICANT: NOEL, MARIANNE
APPLICANT: TAHAR, SOPHIA BEN
APPLICANT: PERRET, JOEL
APPLICANT: BIOSEM
TITLE OF INVENTION: TRANSGENIC PLANTS BELONGING TO THE SPECIES CUCUMIS MELO
FILE REFERENCE: 8076.21USD2
CURRENT APPLICATION NUMBER: US/09/127,742A
CURRENT APPLICATION NUMBER: 08/0309
EARLIER APPLICATION NUMBER: 08/398,209
EARLIER APPLICATION NUMBER: 08/027,563
EARLIER APPLICATION NUMBER: 08/027,563
EARLIER FILING DATE: 1993-03-05
EARLIER FILING DATE: 1993-03-05
EARLIER FILING DATE: 1993-03-05
EARLIER FILING DATE: 1993-03-05
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EARLIER FILING DATE: 1990-08-13
EARLIER APPLICATION NUMBER: FR 89 10848
EARLIER FILING DATE: 1989-08-11
NUMBER OF SEQ ID NOS: 4
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EARLIER FILING DATE: 1995-03-02
EARLIER APPLICATION NUMBER: 08/027,563
EARLIER FILING DATE: 1993-03-05
EARLIER APPLICATION NUMBER: 07/566,255
EARLIER FILING DATE: 1990-08-13
EARLIER APPLICATION NUMBER: FR 89 10848
EARLIER APPLICATION NUMBER: FR 89 10848
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RESULT 13
US-09-252-991A-22748
; Sequence 22748, Application US/09252991A
; Patent No. 6551795
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; LENGTH: 218 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-875-233-10
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                                                                                                                                                                       Query Match
Best Local Similarity
Matches 12; Conserv
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Best Local Similarity
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TELEFAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 10:
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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FILING DATE: 05-MAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 566,255
FILING DATE: 13-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION: NAME: Lisa V. Mueller
                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 312-616-5400
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Reynolds, John F
APPLICANT: Carney, Kim J
TITLE OF INVENTION: Plants Resistant to C Strains
TITLE OF INVENTION: Cucumber Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Chicago
STATE: Illinois
COUNTRY: U.S.A.
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                                                                                                                                   1 MDKSESTSAGRN 12
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6127601
                                                                                                            MDKSESTSAGRN 12
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2 Prudential Plaza, 180 N. Stetson, Suite 4700
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McMaster, J. Russell
Tricoli, David M
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                                                                                                                                                                               85.7%; Score 12; DB 3; Length 218; 100.0%; Pred. No. 3e-05; tive 0; Mismatches 0; Indels
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                                                                                                                                       ; OTHER INFORMATION: Xaa=Arg or Ser US-09-513-999C-7876
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TITLE OF INVENTION: NUCLEIC ACID AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 22748
LENGTH: 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 7876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins. Patent No. 6783961
FILE REFERENCE: 59.US2.REG
                                                           Matches
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Dumas Milne Edwards, J.B. APPLICANT: Duclert, A. APPLICANT: Giordano, J.Y.
                                                                                                                                                                                                                                     LOCATION: 79
OTHER INFORMATION:
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LOCATION: -19..-1
OTHER INFORMATION: SCORE 5.8
OTHER INFORMATION: SEG HLMSLVTSTSASA/SA
                                                                                                                                                                            NAME/KEY: UNSURE LOCATION: 81
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                            NAME/KEY: UNSURE
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7 TSAGRN 12
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vative 0; Mismatches
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TSAGRN 68

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US-09-252-991A-24703
US-09-252-991A 24703, Application US/09252991A

Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: MATC J. Rubenfield et al.
APPLICANT: MATC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: NUCLEIC ACID AND THERAPEUTICS
FILE REFERENCE: 107196,136
CURRENT PELICATION NUMBER: US 09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR PILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1998-07-27
INUMBER OF SEQ ID NOS: 33142
US-09-252-991A-24703
LENGTH: 146
TYPE: PRI
CORGANISM: Pseudomonas acruginosa
US-09-252-991A-24703

Query Match
Best Local Similarity 100.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QU 9 AGRURR 14
Db 131 AGRURR 136

Search completed: December 5, 2004, 08:17:48

Search completed: December 5, 2004, 08:17:48
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RESULT 1 JQ1253 coat protein - cucumber mosaic virus (strain FC) C;Species: cucumber mosaic virus, CMV C;Date: 31 Mar-1992 #sequence_revision 05-Aug-1994 #tex C;Accession: JQ1253 R;Shintaku, M. J. Gen. Virol. 72, 2587-2589, 1991 A;Title: Coat protein gene sequences of two cucumber mo A;Title: Coat protein gene sequences of two cucumber mo A;Reference number: JQ1253; MJID:92013983; PMID:1919534	ALIGNMENTS	45 5 35.7 26 2 804376	42.9 3562 2	6 42.9 2693 2	6 42.9 2570 2	6 42.9 2160 2	6 42.9 1750	6 42.9 1275 2	6 42.9 1143 2	.9 1006 2	6 42.9 742 2	6 42.9 663 1	6 42.9 662 2	6 42.9 662 2	6 42.9 561	6 42.9	6 42.9 497	
rain FC) )5-Aug-1994 #text_change 09-Jul-2004 two cucumber mosaic virus strains reveal a s	TTS	hypothetical prote	chondroitin sulfat	IP3 receptor, XIP3	a				hypothetical prote	probable tail prot	probable kinase/es	gelatinase A (EC 3	gelatinase A (EC 3	gelatinase A (EC 3	65K lower matrix p	65K lower matrix p	hypothetical proté	

## A;Title: Coat protein gene sequences of two cucumber mosaic virus strains reveal A;Reference number: JQ1253; MUID:92013983; PMID:1919534 A;Accession: JQ1254 A;Molecule type: genomic RNA A;Residues: 1-218 <SH1> RESULT JQ1254 R.Tsunasawa, S.; Narita, K. J. Blochem. 92, 607-613, 1982 A;Title: Micro-identification of amino-terminal acetylamino A;Reference number: A61297; MUID:83056735; PMID:6754709 A;Contents: annotation; acetylation C;Accession: JQ1254 R;Shintaku, M. Gen. Virol. 72, 2587-2589, 1991 J. Gen. Virol. 72, 2587-2589, 1991 coat protein - cucumber mosaic virus (strain c/Species: cucumber mosaic virus, CMV C/Date: 31-Mar-1992 #sequence\_revision 05-Aug C/Accession: JQ1254 C;Superfamily: cucumber mosaic virus coat protein C;Keywords: acetylated amino end; coat protein F;1/Modified site: acetylated amino end (Met) #st: A;Cross-references: UNIPROT:Q00259; GB:D10544; NID:g222024; PIDN:BAA01403.1; PID:g222025 A;Note: comparison of the amino acid sequence of this protein to that of strain P6 shows R;Tsunasawa, S.; Narita, K. A;Accession: JQ1253 A;Molecule type: genomic A;Residues: 1-218 <SHI> 밁 á Matches Query Match N Local This cucumber mosaic virus strain induces a yellow chlorosis 14; Similarity MDKSESTSAGRNRR 14 MDKSESTSAGRNRR Conservative RNA 100.0%; 14 0 Score 14; 1 Pred. No. 8 Mismatches 05-Aug-1994 DB 1; of this protein to that of strain FC shows #status experimental .9e-08; #text\_change 09-Jul-2004 Length 218; Indels acids in 0, proteins. Gaps 0 a single single

R;Tsunasawa, S.; Narita, K. J. Biochem. 92, 607-613, 1982 A;Title: Micro-identification

of amino-terminal acetylamino acids in

proteins

A;Cross-references: UNIPROT:Q00261; (A;Note: comparison of the amino acid

GB:D10545; 1 sequence o

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R; Fbunasawa, S.; Narita, N.
J. Biochem. 92, 607-613, 1982
A; Title: Micro-identification of amino-terminal acetylamino acids in proteins.
A; Reference number: A61297; MUID:83056735; PMID:6754709
N. Contents: annotation; acetylation
                                                                                                                                                                                                  A;Title: Comparative studies on A;Reference number: JA0096
A;Accession: JA0097
A;Molecule
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A;Contents: annotation; acetylation
C;Comment: This cucumber mosaic virus strain induces a yellow chlorosis.
C;Superfamily: cucumber mosaic virus coat protein
C;Keywords: acetylated amino end; coat protein
C;Klywords: acetylated amino end (Met) #status experimental
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A;Residues: 216-218 <4RA>>
A;Residues: 216-218 <4RA>>
G;TSunasawa, S; Narita, K.
J. Blochem. 92, 607-613, 1982
A;Title: Micro-identification of amino-terminal acetylamino acids
A;Reference number: A61297; MUID:83056735; PMID:6754709
A;Contents: annotation; acetylation
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A;Title: Complete nucleotide sequence of RNA 3 from cucumber A;Reference number: JS0089; MUID:89279231; PMID:2732698
                                                                                                                                                A;Molecule type: mRNA
A;Residues: 1-218 <NIT>
A;Cross-references: UNIPROT:P18027
                                                                                                                                                                                                                                                                                                                                                                                 coat protein - cucumber mosaic virus (strain Y)
C;Species: cucumber mosaic virus, CMV
A;Note: host Nicotiana tabacum cv. Xanthi nc (tobacco)
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C; Superfamily: cucumber mc
C; Keywords: acetylated ami
F; 1/Modified site: acetyla
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Ann. Phytopathol. Soc. Jpn. 54, 516-522, 1989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Superfamily: cucumber mosaic virus coat protein;
Keywords: acetylated amino end; coat protein;
1/Modified site: acetylated amino end (Met) #st;
                                                                                                                                                                                                                                                                                                                                         Accession: JA0097
                                                                                                                                                                                                                                                                                                                                                              Date: 30-Jun-1990 #sequence_revision 30-Jun-1990
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A;Residues: 1-218 <HYO>
A;Cross-references: UNIPROT:Q66154; EMBL:X77855;
C;Superfamily: cucumber mosaic virus coat proteir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: UNIPROT:Q8JPX2; UNIPROT:Q9DJX2; UNIPROT:Q91PK5; UNIPROT:Q9DJX3; PROT:Q8JPX0; UNIPROT:Q9JPX6; UNIPROT:Q9JPX6; UNIPROT:Q8JPX4; UNIPROT:Q8JPX4; UNIPROT:Q9JPX6; UNIPROT:Q9JPX6; UNIPROT:Q9DWW2; UNIPROT:Q9JWW2; UNIPROT:Q9DWW2 A;Experimental source: strain CS A;Note: the authors translated the codon CGG for residue 83 as Gly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       coat protein - cucumber mosaic virus (strain CS)
C;Species: cucumber mosaic virus, CMV
C;Date: 21-Jan-1997 #sequence_revision 21-Jan-1997
C;Accession: JC6075
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Map position: segment RNA3
c;Superfamily: cucumber mosaic virus cc
C;Keywords: acetylated amino end; coat
F;1/Modified site: acetylated amino end
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                              Ş
                                                                                                                                                                                  A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-218 <HYO>
                                                                                                                                                                                                                                                             submitted to the EMBL Data Library, February 1994 A;Description: Nucleotide sequence of cDNA from RA;Reference number: S42098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Huaping, L.; Hu, J.S.; Barry, K.; Hweichung, F. Chinese J. Virol. 12, 235-242, 1996
A;Title: Coat protein gene sequence analysis of ta, Reference number: JC6073
                                                                                                                                                                                                                                                                                                                 R; Hyon, S.; Park,
                                                                                                                                                                                                                                                                                                                                     C; Accession:
                                                                                                                                                                                                                                                                                                                                                    C;Species: cucumber mosaic virus, CMV
C;Date: 06-Jan-1995 #sequence_revision
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C; Superfamily: cucumber mosaic virus
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A; Residues: 1-218 < HUA>
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                                                                         Matches
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                                  1 MDKSESTSAGRNRR 14
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                                                                       Conservative
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                                                                   100.0%; Score 14; DB 2; I
100.0%; Pred. No. 8.9e-08;
Live 0; Mismatches 0;
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Pred. No.
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IPROT:Q9DC
9DWW2; UNI
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C;Species: cucumbe
C;Date: 13-Jan-199
C;Accession: S5803
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S58039
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A;Cross-references: UNIPROT:Q83257;
A;Experimental source: strain MM
A;Cross-references: UNIPROT:Q9DJW8; UNIPROT:Q9YJR7; UNIPROT:Q8JPX1; UNIPROT:Q9E2V6; UNIPROT:Q91PP6; UNIPROT:Q9JPY4; UNIPROT:Q66157; UNIPROT:Q66130; UNIPROT:Q8JPX4; UNIPROT:Q9GDW2; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9JPX4; UNIPROT:Q9
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A;Title: Coat protein gene sequence analysis o. A;Reference number: JC6073
                                                                                                                                                                                                                                                                                                                                                             C;Species: cucumber mosaic virus, CMV
C;Date: 21-Jan-1997 #sequence_revision 21-Jan-1997
C;Accession: JC6073
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Chinese J. Virol. 12, 235-242, 1996
A;Title: Coat protein gene sequence analysis of three cucumber
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A;Residues: 1-218 <HAQ>
A;Cross-references: UNIPROT:Q66157; EMBL:X89652; NID:g902619; PIDN:CAA61802.1; PID:g902C;Superfamily: cucumber mosaic virus coat protein
                                                                                                                                 A; Molecule type: mRNA
A; Residues: 1-218 < HUI
                                                                                                                                                                                                       A;Accession: JC6073
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A; Accession: JC6074
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Species: cucumber mosaic virus, CMV
Date: 21-Jan-1997 #sequence_revision 21-Jan-1997
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Best Local
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                                                                                                                             1-218 <HUA>
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Pred. No.
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Pred. No. 8.9e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UNIPROT: 055251; UNIPROT: Q9WAB3;
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A;Gene: cp
C;Superfamil
C;Keywords:
                                                                                                                              R;Quemada, H.; Kearney, C.; Gonsalves, D.; Slightom, J.L. J. Gen. Virol. 70, 1065-1073, 1989
A;Title: Nucleotide sequences of the coat protein genes a A;Reference number: JU0087; MUID:89279284; PMID:2732712
A;Accession: JA0136
                                                                                                                                                                                                                          coat protein - cucumber mosaic virus (strain C)
C;Species: cucumber mosaic virus, CMV
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change
C;Accession: JA0136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Experimental source: strain I17F
A;Note: this is a revision to the sequence from reference S06932
R;Noel, M.J.T.; Tahar, S.B.
Nucleic Acids Res. 17, 10492, 1989
A;Title: Nucleotide sequence of the coat protein gene and flankin A;Reference number: S06932; MUID:90098885; PMID:2602158
A;Accession: S06932
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R;Noel, M.J.T.; Tahar, S.B.
Nucleic Acids Res. 18, 1332, 1990
A;Title: Corrigendum. Nucleotide sequence of the coat protein A;Reference number: S09663; MUID:90206832; PMID:2320436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Variety: strain I17F
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change
C;Accession: S09663; S06932
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S09663
R;Tsunasawa, S.; Narita, K.
J. Biochem. 92, 607-613, 1982
A;Tille: Micro-identification of amino-terminal acetylamino
A;Reference number: A61297; MUID:83056735; PMID:6754709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Molecule type: mRNA
A;Residues: 1-6,'W',8-44,'W',46-51,'W',53-56,'W',58-68,'W',70,'W',72-73,'W',75-96,'W',98
A;Cross-references: EMBL:X16386
                                                                                             A; Molecule type: genomic RNA
A; Residues: 1-218 < QUE>
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A;Note: this sequence has been revised in reference
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A; Residues: 1-218 < NOE>
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                                                                         A;Cross-references: UNIPROT:P21368; GB:D00462; NID:g222041;
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Pred. No. 8.
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Pred. No. 8.9e-08
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annotation;

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hypothetical 11.5K protein (clone GV-B) - garlic virus B
N;Alternate names hypothetical protein II
C;Species: garlic virus B
C;Date: 30-Sep-1993 #sequence_revision 20-Aug-1994 #text_change 09-Jul-2004
C;Accession: JQ2176; S27912
R;Sumi, S.; Tsuneyoshi, T.; Furutani, H.
J. Gen. Virol. 74, 1879-1885, 1993
A;Reference number: JQ2171; MUID:93389442; PMID:8376963
A;Accession: JQ2176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Accession: D71392
R;McGarvey, P.; Tousignant, M.; Geletka, L.; Cellini, F.; Kaper, J.M. J. Gen. Virol. 76, 2257-2270, 1995
A;Title: The complete sequence of a cucumber mosaic virus from Ixora A;Reference number: A71392; MUID:96005047; PMID:7561763
A;Accession: D71392
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C;Superfamily: cucumber mensic virus coat protein
C;Keywords: acetylated amino end; coat protein; glycoprotein
F;1/Modified site: acetylated amino end (Met) #status experimental
F;43/Binding site: carbohydrate (Asn) (covalent) #status predicted
                                                                                                                 R;Sumi, S.I.
submitted to the EMBL Data
A;Reference number: S27908
A;Accession: S27912
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C;Comment: The CMV genome consists of three species of single-stranded, capped, positive
C;Comment: The Ixora strain in unusual in that it does not replicate several well charace
ptoms found when replicated by other CMV strains.
C;Genetics:
                                                                                                                                                                                                          A;Molecule type: mRNA
A;Residues: 1-108 <SUM>
A;Cross-references: UNIPROT:Q67695; GB:AB010301;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             coat protein - cucumber mosaic virus (strain Ixora)
N;Alternate names: capsid protein
C;Species: cucumber mosaic virus, CMV
C;Date: 03-Aug-1998 #sequence_revision 03-Aug-1998 #text_change 09-Jul-2004
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A;Cross-references: EMBL:D11158; NID:g221432; PIDN:BAA61814.1; C<sub>A</sub>Superfamily: barley stripe mosaic virus 14K protein
                                             A; Molecule type: DNA
A; Residues: 1-108 <SU2>
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                                                                                           Status: preliminary
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Best Local
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100.0%; Pr
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100.0%; Pred. No.
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                                                                                                                                                                 July 1992
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                                                                                                                                                                                                             GB:D11158; NID:g2826152; PIDN:BAA61814
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A;Molecule type: DNA
A;Residues: 1-133 <GRA>
A;Cross-references: UNIPROT: (084440) EMBL: U42580; NID: 94028896;
A;Experimental source: specific host Chlorella strain NC64A
C;Genetics:
                                                                                                                                                                                                                                                                                                 C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 09-Jul-2004 C;Accession: T47771  
C;Accession: G; Fartmann, B.; Dauner, D.; Sterr, W.; Holland, R.; Weichselgartner, M.; N submitted to the Protein Sequence Database, February 2000 A;Reference number: Z24475  
A;Accession: T47771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hypothetical protein al19L - Chlorella virus PBCV-1
C;Species: Chlorella virus PBCV-1
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
C;Accession: T17609
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Best Local Similarity
Tatches 6; Conserve
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 09-Jul-2004
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A;Reference number: Z18806
A;Accession: T17609
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A; Note: F24I3.190
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A;Experimental source: cultivar Columbia; BAC clo
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A; Residues: 1-136 < NYA>
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Best Local (
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SESTSA 11
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100.0%; Pred. No. 18;
tive 0; Mismatches
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100.0%; Pred. No.
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